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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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 A Geneseq 21:*
1: geneseqp1990
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37
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              geneseqp1908:*
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34.688 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

20 25 67.6 21 25 67.6 22 25 67.6 23 25 67.6	25 67. 25 67. 25 67.	25 67. 25 67.	25 67. 25 67.	25 67.	25 67.		19 25 67.6		25 67.	17 25 67.6	25		26 70	13 28 75.7		36 97		97	36 97	36 97	6 36 97.3	36 97	36 97.	36 97.	2 36 97.3	. 36 97.	No. Score Match Length	Query
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The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the

A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.

Claim 1; Page 32; 47pp; English.

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Human can	Breast ca	Breast ca	Streptomy	SARS vira	Severe ac	SARS coro	Human can	Y. pestis	SARS coro											

ALIGNMENTS

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RESULT 1
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ID AAW99
XX AAW99
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                                                                                                                                                                                              The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency
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AAW99197 standard; peptide; 9
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RESULT 4
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Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-and T-cells
Typing minor histocompatibility antigen HA-1 -
                                                  WPI; 1999-142960/12
                                                                                                     Goulmy
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02-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HA-1 H-allele sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-142855/12
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nes 9; Conserv
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by amplifying
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The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as ever detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                    Typing minor histocompatibility antigen HA-1 - identifying HA-1 R and HA-1 H alleles of HA-1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                          Claim 13; Fig
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of genetic aberrances.
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98EP-00870125.
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88.9%;
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                                                                                                   The invention relates to a novel method for providing epitopes of allelic convariants of antigenic proteins from specific species based on single concleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein/peptide or its converged to the protein of the invention, converged by DNA sequence having SNP, creating variant epitopes, converged by DNA sequence having SNP, creating variant epitopes, converged by DNA sequence having SNP, creating variant epitopes, converged by DNA sequence having SNP, creating variant epitopes, converged by DNA sequence having SNP, creating variant epitopes, converged by applying the method is converged by applying the method is converged by applying the method for several protein from the converged by applying the method for several protein from the converged by applying the method for several protein from the converged by applying the sprofile was related to disease, preferably cancer. This is useful for diagnosing a disease in an individual by converged by applying the SNP-related polymorphic profile. A method of the invention converged by the progression, regression or onset of a treated disease. The present converged by the progression, regression or onset of a treated disease. The present converged by the progression in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 82; 119pp; English.
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                                                                       Sequence 9
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                                                                                                                                                                          The invention relates to a novel method for providing epitopes of allelic convariants of antigenic proteins from specific species based on single conclectide polymorphism (SNP), by defining target protein/peptide or its converted to the protein protein dentifying, conclecting allelic peptide/protein variants, expression product or its configuration of the invention has cytostatic activity, and may have a use in a vaccine. The method is concertified by applying the method for several protein from the concertifies where the SNP profile of one or more individuals from a concertified where the SNP profile was related to disease, preferably cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by concertified the SNP-related polymorphic profile. A method of the invention concertified the snP-related polymorphic profile. A method of the invention concertified the snP-related polymorphic profile. A method of the invention concertified the snP-related polymorphic profile. A method of the invention concertified polymorphic profile is determining cancer, preferably leukaemia, and for determining concertified polymorphic profile. A method of the invention concertified polymorphic profile is demanded to disease. The present concertified in the exemplification of the invention.
                                                                      Query Match
Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Strittmatter W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAR-2004
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                                                                                                                                              Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 82; 119pp; English.
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                                                                        Score 36; DB
Pred. No. 2e+0
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RESULT 9
AAW99199
ID AAWS

AAW99199 standard; peptide; 13

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RESULT 10
AAW99198
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Best Local
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                                                       Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
           WO9905174-A1
                                                                                                         DH cell KIAA0223 protein sequence
                                                                                                                                   20-MAY-1999
                                                                                                                                                                                AAW99198 standard; peptide;
                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 31; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-153312/13.
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                                                                                                                                                                                                                                                                                                                                                                       invention
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Pred. No.
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Best Local Similarity
Matches 8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a new peptide (PI) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. PI is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases. The present sequence represents a KIAA0223 sequence given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                 AAW97415 standard; protein; 13
Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
                                                                              Goulmy EAJM,
                                                                                                                                                           23-JUL-1998;
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                                                                                                                                                                                                                                       Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
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                                                                                                        (UYLE-) RIJKSUNIV LEIDEN.
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                                         1999-142855/12.
DB; AAX16081.
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                                                                                                                                                                                                                                                                                                                  polymorphism in HA-1 positive homozygous individuals
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                                                                              Hunt DF,
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Pred. No. 1.8;
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Disclosure;

Page 38;

57pp;

English.

versus-host disease

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RESULT 12
AAW97414
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease; neoplastic haematopoietic cell; KIAA0223 polymorphism.
The present sequence represents the KIAA0223 polymorphism in HA-1 negative homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
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                                                                                                                                                                                                                                                         Immunogenic peptide from minor histocompatibility antigen HA-1 - usef for inducing tolerance to transplants and prevent rejection or graft-
                                                                                                                                                                                                                                                                                                                                                              Goulmy EAJM, Hunt DF,
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                                                                                                                                                                                                   Disclosure; Page 38;
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RESULT 13
ABP71119
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XX ABP71119
XX ABP71119
XX ABP71
XX ABP71
XX ABP71
XX ABP71
XX BTF3
KW BTF3
KW BTF3
KW NOCO
XX Unida
XX E1-M
XX E21-M
XX 21-M
XX 21-M
XX Inhi
PT 21-M
XX Inhi
PT e.g.
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Best Local S
Matches 8
                                                              Query Match
Best Local S
Matches 6
                                                                                                                                                         The invention relates to inhibiting or increasing programmed cell death of a cell. The method involves upregulating or inhibiting, respectively, the expression or activity of basic transcription factor (BTF) or its homologue in the cell. The BTF3 polypeptides and nucleic acids are useful for inhibiting or increasing programmed cell death. They are used for screening for an agent that increases or inhibits programmed cell death or pre-screening for an agent that modulates programmed cell death. The screened agent that increases or inhibits programmed cell death, is used for diagnosing or treating cancer or neurodegenerative disease (e.g. amyotrophic lateral sclerosis, Alzheimer's disease, Parkinson's disease or multiple sclerosis), atherosclerosis, or rheumatoid arthritis. Sequences ABF71106-123 represent CARD regions of various CARD proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BTF3; cell death; apoptosis; basic transcription factor; cytostatic; nootropic; neuroprotective; antiparkinsonian; antiarteriosclerotic; antirheumatic; antiarthritic; gene therapy; CARD; E10.
                                                                                                                                                                                                                                                                                                                                                                                                                   Inhibiting or increasing programmed cell death of a cell, for treating e.g. cancer, comprises upregulating or inhibiting, respectively, the expression or activity of basic transcription factor (BTF)3 or its homolog in the cell.
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3 LVDDLLE
                               2 LXDDLLE
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                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bloss T,
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Pred. No. 1.8;
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Pred. No.
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RESULT 15
AAW73901
ID AAW73
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AC AAW73
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                                                                                                                                                                                                                                                                                     CC This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein). a polymerase protein L and CC SH (also known as the 1A protein). Specifically, it refers to a CC composition comprising an above peptide mixed with a pharmaceutical excipient or an RSV immunogenic composition comprising a recombinant CC excipient or an RSV immunogenic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC (MHC) class I restricted T-cell stimulating peptide pitope of RSV. The CC RSV peptide is useful for preparing a diagnostic composition or an RSV (prophylactic or therapeutic) vaccine composition for a DNA-based CC (immunisation, or for preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the polypeptide in an immune response provoking an activity. This peptide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                      Query Match
Best Local S
Matches 4
                AAW73901;
                                          AAW73901 standard; peptide;
                                                                                                                                                                                                                                                   Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-758334/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human respiratory syncytial virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human respiratory syncytial virus; cytotoxic T lymphocyte; CTL; MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human RSV L high affinity binding peptide SeqID 919
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Matches
                                                                                                                   Query Match
                                                                                                                                           Sequence 10 AA;
                                                                                                                                                               This sequence is a fragment of the Bacteroides forsythus protease of the invention. The protein has protease activity and haemolytic activity. The DNA and the polypeptide are useful for diagnosis, treatment and research of periodontosis. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                        Disclosure; Page 13; 13pp; Japanese.
                                                                                                                                                                                                                                             New protease gene derived from Bacteroides forsythus - useful for diagnosis and treatment of periodontosis.
                                                                                                                                                                                                                                                                                                                                                                                                    JP11009284-A.
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12-APR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Protease; haemolytic activity; diagnosis; periodontosis.
                                                                                                                                                                                                                                                                                                                                                                                                                         Tannerella forsythensis.
                                                                                                                                                                                                                                                                                                         (SUNZ ) SUNSTAR CHEM IND CO LTD
                                                                                            Local Similarity nes 4; Conserv
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Pred. No.
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OM protein protein search, using sw model Copyright GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.

Title: Perfect score: Sequence: US-10-791-217A-1 37 VLXDDLLEA 9

Run on:

April

6, 2006, 16:57:26 ; Search time 19 Seconds (without alignments)
45.576 Million cell updates/sec

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283416 segs, 96216763 residues

Total number of hits satisfying chosen parameters:

2523

Minimum DB Maximum DB seq length: 0 seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

PIR 80:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

ALIGNMENTS

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RESULT 2 C26997

Unspecific monooxygenase (EC 1.14.14.1) isozyme E, phenobarbital-inducible, hepatic - ra
W,Alternate names: cytochrome P450
C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 31-Dec-2004
C;Accession: C26997
R;Graves, P.E.; Kaminsky, L.S.; Halpert, J.
Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional and structural multiplicity of pregnenolone-16-alpha-ca
A;Feference number: A26997; MUID:88000604; PMID:3651420
A;Accession: C26997
A;Molecule type: protein
A;Residues: 1-10 <GRA>
A;Cross-references: UNIPROT:006884; UNIPARC:UPI0000174DA8
C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase; oxidoreductase; transmembrane

ঠ Matches Query Match Best Local (Local Similarity 5 DLLEA 9 4 Conservative 48.6%; 0, Score 18; DB 2; Pred. No. 9e+02; 0; Mismatches 1; Length 10; Indels 0, Gaps o;

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DLLSA

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R;Ohishi, N.; Imaoka, S.; Suzuki, T.; Funae, Y. Biochim. Biophys. Acta 1188, 227-236, 1993 A;Title: Characterization of two P-450 isozymes placed in the A;Reference number: S39761; MUID:94072607; PMID:8251521 A;Accession: S39762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Rattus norvegicus (Norway rat)
C;Date: 19-May-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: S39762
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S39762
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                                                                                                                                                                        virg protein - Agrobacterium sp. (fragment)
C;Species: Agrobacterium sp.
C;Species: Agrobacterium sp.
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_change 09-Jul-2004
C;Accession: S13224
R;Tamamoto, S:; Aoyama, T.; Takanami, M.; Oka, A.
J. Mol. Biol. 215, 537-547, 1990
A;Title: Binding of the regulatory protein Virg to the phased signal sequences upstream
A;Reference number: S13224; MUID:91039316; PMID:2231718
A;Accession: S13224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: protein A; Residues: 1-15 < GRA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional and structural multiplicity of pregnenolone-16-alpha-ca A;Reference number: A26997; MUID:88000604; PMID:3651420
A;Accession: A26997
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A;Residues: 1-12 <OHI>
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                                                                                                     A;Status: preliminary
A;Molecule type: protein
A;Residus: 1-10 «TAM»
A;Cross-references: UNIPROT:Q7M0P7; UNIPARC:UPI000017A9C4
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Pred. No. 1.5e+03;
1; Mismatches 1.
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1.1e+03,
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   A; Molecule type: protein
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potassium channel protein Slo G3 - fruit fly (Drosophila melanogaster) (fragment) C;Species: Drosophila melanogaster C;Species: Drosophila melanogaster C;Apecies: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Feb-1997 C;Accession: PS0443 R;Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta, A Reprosophila Reprosophi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T-cell receptor beta chain V-D-J region (isolate 10) - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997 C;Accession: PH0924
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A;Experimental source: concanavalin A-activated lymphoblast C;Keywords: T-cell receptor
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A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental A;Reference number: PH0891; MUID:92078857; PMID:1836012
                                                                                                                                      chlorophyll a/b-binding protein Lhcb5 - spinach (fragment)
N;Alternate names: light-harvesting complex LHCIIc protein
C;Species: Spinacia oleracea (spinach)
C;Date: 19-Mar-1997 #sequence_revision 23-Apr-1999 #text_C
C;Accession: S50900
C;Accession: S50900
R;Walters, R.G.; Ruban, A.V.; Horton, P.
B;Walters, R.G.; Ruban, A.V.; Horton, P.
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A;Molecule type: mRNA
A;Residues: 1-13 <ADE>
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A; Residues: 1-11 <GOL>
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                             Eur. J. Biochem. 226, 1063-1069, 1994
A; Title: Higher plant light-harvesting complexes LHCIIa
A; Reference number: S50900; MUID:95112835; PMID:7813461
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C;Keywords: alternative splicing; ion c
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A; Accession: S50900
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Pred. No. 1.6e+03
1; Mismatches
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PMID:1497890
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S62675
collagen type I - mouse (fragment)
c;Species: Mus musculus (house mouse)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S62675
R;Mizuno, M.; Kitafima, T.; Tomita, M.; Kuboki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                PS I complex subunit 8 - cucumber (fragment)
(;Species: Cucumis sativus (cucumber)
(;Species: Cucumis sativus (cucumber)
(;Date: 05-Jan-1996 #sequence_revision 05-Jan-1996 #text_change 09-Jul-2004
(;Accession: E56819
R;Iwasaki, Y; Ishikawa, H.; Hibino, T.; Takabe, T.
Biochim. Biophys. Acta 1059, 141-148, 1991
A;Title: Characterization of genes that encode subunits of cucumber PS I con A;Reference number: A56819; MUID:91355209; PMID:1883835
A;Accession: E56819
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c;Species: Persea americana (avocado)
C;Date: 24-Uan-1992 #sequence_revision 24-Jan-1992 #text_change 03-Jun-2002
C;Accession: C39170
R;Shanklin, J.; Somerville, C.
R;Shanklin, J.; Somerville, C.
Proc. Natl. Acad. Sci. U.S.A. 88, 2510-2514, 1991
A;Title: Stearoyl-acyl-carrier-protein desaturase from higher plants is structurally unz
A;Reference number: A39170; MUID:91172837; PMID:2006187
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A;Cross-references: UNIPARC:UPI0000178182
C;Superfamily: chlorophyll a/b-binding protein
C;Keywords: chlorophyll; chloroplast; light-harvesting complex; photosynthesis; photosys
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A;Note: sequence extracted from NCBI backbone (NCBIP:5860
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A;Molecule type: protein
A;Residues: 1-14 <SHA>
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A;Accession: C39170
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Pred. No. 2.3e+03
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A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: CAN>
A;Residues: 1-8e (ANA>
A;Residues: CANA>
A;Cross-references: UNIPROT:P95412; UNIPARC:UPI000017A96A; DDBJ:D50473; NID:g1217594
A;Note: this ORF is not annotated in GenBank entry PSENIRC, release 113.0
C;Superfamily: Pseudomonas stutzeri nirD protein
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A;Title: Sequencing and characterization of the downstream y for biosynthesis of heme dl.
                                                                                                                                                                                                                                                                                                                                                                                                                 hypothetical protein 8 [imported] - Pseudomonas aeruginosa (fragment) c;Species: Pseudomonas aeruginosa (c;Species: Pseudomonas aeruginosa (c;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 31-Dec-2004 c;Accession: PC4131 R;Kawasaki, S.; Arai, H.; Igarashi, Y.; Kodama, T.
                                                                                                                                                                                                                                                                                                           A;Reference number: JC4552; MUID:96144254; PMID:8566817 A;Accession: PC4131
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A;Title: Hucolin, a new corticosteroid-binding protein from A;Reference number: $68004; MUID:96087107; PMID:7498469
A;Accession: $68004
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 <EDG>
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A,Title: The osteoblastic MC3T3-E1 cells synthesized C-terminal propeptide of type I A,Reference number: 862675; MUID:97386332; PMID:9244181
A;Accession: 862675
A;Accession: S62675
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-15 <MIZ>
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C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
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carnitine medium/long chain acyltransferase (EC 2.3.1.-) - rat (fragment)
N;Alternate names: endoplasmic reticulum protein ERp61; glucose regulated protein GRP58;
C;Species: Rattus norvegicus (Norway rat)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-May-1999
C;Accession: A61354
R;Murthy, M.S.R.; Pande, S.V.
Mol. Cell. Biochem. 122, 133-138, 1993
A;Title: Carnitine medium/long chain acyltransferase of microsomes seems to be the previ A;Reference number: A61354; MUID:94049728; PMID:8232244
A;Accession: A61354
A;Ccession: A61354
A;Accession: A61354
A;Accession: A61354
A;Residues: 1-10 cMUR>
A;Cross-references: UNIPARC:UPI000017C8DE
C;Keywords: acyltransferase
                                                                                                                                                                                                                                                                                                photosystem I 9K chain - spinach (fragment)
C;Species: Spinacia oleracea (spinach)
C;Ante: 19-Mar-1997 #sequence_revision 13-Mar-1998 #text_change 09-Jul-2004
C;Antesion: S14316
R;Ikeuchi, M.; Inoue, Y.
FEBS Lett. 280, 332-334, 1991
A;Title: Two new components of 9 and 14 kDa from spinach photosystem I complex.
A;Reference number: S14316; MUID:91192162; pMID:2013332
A;Accession: S14316
A;Accession: S14316
A;Accession: S14316
A;Kolecule type: protein
A;Residues: 1-13 < IKE>
A;Cross-references: UNIPROT:Q7M1J1; UNIPARC:UPI000017AF28
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J. Biol. Chem. 269:12263-12268(1994).
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RESULT 5
Q9UC60 HUMAN
Q9UC60 HUMAN PRELIMINARY;
AC Q9UC60;
DT 01-MAY-2000 (TrEMBLrel. 13, La
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DT 01-JUN-2003 (TrEMBLrel. 24, La
DT 01-JUN-2003 (TrEMBLrel. 24, La
DE N-acetylmuramyl-L-alanine amid
OS Homo sapiens (Human).

CE Lukaryota; Metazoa; Chordata;
OC Homo.
OX NCBI_TaxID=9606;
RN [1]
RP PROTEIN SEQUENCE
RY [1]
RP PROTEIN SEQUENCE
RY "Characterization of human sea
RT "Characterization of human sea
Purified by affinity chromatog
RY GO; GO:0001519; P:peptide amid
SQ SEQUENCE 15 AA; 1600 MW;
Q9S5D6 CHRVI
ID Q9S5D6 CHRVI
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J. Bacteriol. 173:1367-1375(1991).
EMBL; M63619; AAA24556.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDITINE-95397215; PubMed=7663175; DOI=10.1006/prep.1995.1049; DD Pauw P., Neyt C., Vanderwinkel E., Wattiez R., Falmagne P.; "Characterization of human serum N-acetylmuramyl-L-alanine amic purified by affinity chromatography."; protein Expr. Purif. 6:371-378(1995). GO:0008745; F:N-acetylmuramoyl-L-alanine amidase activity; GO:GO:0001519; P:peptide amidation; NAS. SEQUENCE 15 AA; 1600 MW; 9016B00FFF9E780A CRC64;
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                     Liebergesell M., Schmidt B., Steinbuchel "Isolation and identification of granule-for poly(3-hydroxyalkanoic acid) biosynth D.";
                                                                                                                                                                                                                                                               Bacteria; Proteobacteria; Gamm
Chromatiaceae; Allochromatium.
                                                                                                                                                                                                                                                                                                                 (Fragment).
Chromatium vinosum.
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     SEQUENCE
     FEMS Microbiol.
SEQUENCE 15 AJ
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MEDLINE=93146381; PubMed=1490603; DOI=10.1016/0378-1097(92)90031-I;
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Conservative
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       Lett. 78:227-232(1992).
A; 1874 MW; 165FA9A16BCA2A9D
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MEDLINE=92291078;
Subramaniam V.N.,
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                                                                                                                                                                        Golgi-enriched membranes.";
J. Biol. Chem. 267:12016-12021(1992).
                                                                                                                                                                                                                                                                                                                                                                     Rattus norvegicus (Rat).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna;
Muridae; Murinae; Rattus.
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2000 (TrEMBLrel. 14,
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Q718N9_9PARA PRELIMINARY;
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Della-Porta A.J.;
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Paramyxoviridae; Paramyxovirinae; Avulavirus.
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bin Mohd Yusoff A.R.,
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RESULT 9
Q718K6 9PARA
Q718K6 9PARA PRELIMINARY;
AC
Q718K6;
DT
05-JUL-2004 (TYEMBLrel. 27,
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05-JUL-2004 (TYEMBLrel. 27,
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DT
Emagglutinin-neuraminidase
OS
Newcastle disease virus.
OC
Viruses; ssRNA negative-str;
OC
Paramyxoviridae; paramyxovi.
OX
NCBI TaxID=11176;
RN
(I)1
RP
NUCLEOTIDE SEQUENCE.
RA
GOULd A.R., Hansson E., Sel.
RA
GOULd A.R., Hansson E., Sel.
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Could A.R., Hansson E., Sel.
RA
Della-Porta A.J.;
RT
"Newcastle disease virus fur
motifs as markers for viral
RL
Avian Pathol. 32:361-373 (20)
DR
EMBL; AP542917; AAQ11642.1;
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SEQUENCE 11 AA; 1313 MW;
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Best Local S
Matches 3
HAWAP; MF 00455; -; 1.

HAWAP; MF 00455; -; 1.

InterPro; IPR001998; Xylose isom.

PROSITE; PS00172; XYLOSE ISOMERASE 1; PARTIAL.

PROSITE; PS00173; XYLOSE ISOMERASE 2; PARTIAL.

PROSITE; PS00173; XYLOSE ISOMERASE 2; PARTIAL.

Carbohydrate metabolism; Direct protein sequencing; Isomeras

Carbohydrate metabolism; Pentose shunt; Xylose metabolism.
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10-MAY-2005 (Rel.
Xylose isomerase
                                                                                                                                                                                                                                                                                                                                   mapping.";
Biochem. J
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P14405;
                                                                                                                                                                                                                                                                                                                                                               MEDLINE=90104230; PubMed=2604694; Vangrysperre W., Ampe C., Kersters-Hilderson H., Tempst P.; Wangrysperre W., Ampe C., Kersters-Hilderson H., Tempst P.; Wingle active-site histidine in D-xylose isomerase from St violaceoruber. Identification by chemical derivatization an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptomyces violaceoruber.
Bacteria; Actinobacteria; A
                                                                                                                                                                            This Swiss-Prot entry is copyright. It is produced through a obstween the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                        -!- SUBUNIT: Homotetramer.
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the xylose isomerase family.
                                                                                                                                                                                                                                                                                               Biochem. J. 263:195-199(1989).
-!- CATALYTIC ACTIVITY: D-xylose = D-
-!- COFACTOR: Binds 2 magnesium ions
                                                                                                                                                                                                                                                                                                                                                                                                                                    PROTEIN SEQUENCE.
STRAIN=LMG 7183;
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Avian Pathol. 32:361-373(2003).
EMBL; AF542917; AAQ11642.1; -; Genomic_DNA.
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                                                                                                                                                          use as long as its content is in no way modified
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13, Last sequence update)
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Pred. No. 9.7e+03;
1; Mismatches 1;
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and this statement
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Best Local
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Q9TQY4;
Q9TQY4;
01-MAY-2000 (TrEMBLrel. 13, Cr
01-MAY-2000 (TrEMBLrel. 21, La
01-JUN-2002 (TrEMBLrel. 21, La
Glycoprotein H-B N-TERMINAL, C
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01-MAY-2000 (TrEMBLrel. 12, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
NADH:ubiquinone oxidoreductase (Complex I) iron-sulfur
fraction 20 kDa polypeptide peptide T-8 (Fragment).
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                       MEDLINE=91340701; PubMed=1874725;
Christie D.L., Batchelor D.C., Palmer D.J.
"Identification of kex2-related proteases
partial amino acid sequence analysis.";
J. Biol. Chem. 266:15679-15683(1991).
                                                                                                                                                               Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; !
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;
Pecora; Bovidae; Bovinae; Bos.
NCBI_TaxID=9913;
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Bos taurus (Bovine).
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J. Biochem. 110:575-582(1991).
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Masui R., Wakabayashi S., Matsubara H.
"The amino acid sequence of the 9 kDa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;
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Pred. 1
1; Mis
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Pred. No. 1.1e
1; Mismatches
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RESULT 14
Q5MK52 9CLOS I
D K55MK52 9CLOS I
AC Q5MK52;
DT 01-FEB-2005 (7)
DT 01-FEB-2005 (7)
DT 01-FEB-2005 (7)
DE RdRp (Fragment
OS Citrus triste
OC Viruses; ssRNJ
OC Closterovirus
OX NCBI TAXID=121
RN [1]
RP NUCLEOTIDE SEC
RP NUCLEOTIDE SEC
RP STRAIM=8358;
RA Hilf M.B., Man
RT "POL and K17 "
RL Submitted (SEI
DR EMBL; AY5630;
FT NON TER
SQ SEQUENCE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QSMK49 9
AC Q5MK49 9
AC Q5MK49 7
DT 01-FEB-2
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DT 01-FEB-2
DT 01-FEB-2
DT 01-FEB-2
CO CITUBE FT OC CLOSTERC
OC CLOSTERC
OC VITUSES
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01-FEB-2005
01-FEB-2005
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"POL and K17 markers from Beltsville CTV isolates.";
Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databa
EMBL; AY756309; AAW22818.1; -; Genomic_RNA.
                                                                                                                                                                                  STRAIN-B358;
Hilf M.E., Mavrodieva V.A., Garnsey S.M.;
"POL and K17 markers from Beltsville CTV isola
Submitted (SEP-2004) to the EMBL/GenBank/DDBJ
EMBL; AY756307; AAW22815.1; -; Genomic_KNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=B359;
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Viruses; ssRNA positive-strand
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                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=12162;
                                                                                                                                                                                                                                                                                                                                                       Closterovirus.
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Last annotation updat
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Pred. No.
                                                                             Pred. No. 1.1
1; Mismatches
                                                                                               Score 18;
Pred. No.
                                                                                                                                                          CAA485D77F187DC6
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Page 5
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RESULT 15
Q5MK66 9CLOS PRELIMINARY; PRT; 12 AA.

AC Q5MK66;
DT 01-FEB-2005 (TEMBLrel. 29, Created)
DT 01-FEB-2005 (TEMBLrel. 29, Last sequence update)
DT 01-FEB-2005 (TEMBLrel. 29, Last sequence update)
DT 01-FEB-2005 (TEMBLrel. 29, Last sequence update)
DE RdEp (Fragment).
OS Citrus tristera virus.
OC Viruses; SERNA positive-strand viruses, no DNA stage; Closteroviridae;
OC Closterovirus.
OC Closteroviru
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THIS PAGE BLANK (USPTO)

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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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         91:
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seq length: 15
         protein search, using sw model
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Match
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/Backfiles1.pep:*
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37
1 VLXDDLLEA 9
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         GenCore version 5.1.7
(c) 1993 - 2006 Biocceleration
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                  US-09-269-250E-18
US-09-269-250E-29
US-09-489-760-1
US-09-489-760-2
US-09-489-760-2
US-09-489-760-2
US-09-489-760-1
US-09-269-250E-26
US-09-269-250E-26
US-09-269-250E-38
US-09-489-760-16
US-09-489-760-11
US-09-489-760-10
US-09-489-760-10
US-08-940-095-104
US-08-940-095-104
US-08-940-105-104
US-08-940-105-104
US-09-453-803-104
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Sequence
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20, Appl
20,
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US-09-269-250E-20 (US-09-269-250E-20 (US-09-269-260E-20 (US-09-269-260E-20 (US-09-269-260E-20 (US-09-260E-20 (US-09-	Query Match 97.3%; Score 36; DB 2; Length 9; Best Local Similarity 88.9%; Pred. No. 4.6e+05; Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0; Qy 1 VLXDDLLEA 9 Db 1 VLRDDLLEA 9	RESULT 1 US-09-269-250E-18 Sequence 18, Application US/09269250E Patent No. 6830883 GENERAL INFORMATION: APPLICANT: Goulmy, Elsa ITILE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 PILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: US/09/269,250E CURRENT FILING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 18 LENGTH: 9 TYPE: PRT ORGANISM: Artificial Sequence PEATURE: OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-09-269-250E-18	28 22 59.5 9 2 US-09-744-549-15 29 22 59.5 9 2 US-10-394-980-421 30 22 59.5 9 2 US-10-394-980-421 31 22 59.5 11 US-08-726-136-13 32 25 59.5 11 2 US-09-103-434-13 32 25 59.5 11 2 US-09-103-434-13 33 22 59.5 11 2 US-09-687-594-13 34 22 59.5 12 2 US-09-687-594-13 35 22 59.5 14 1 US-09-366-276-9 36 22 59.5 15 2 US-09-069-8277A-132 37 22 59.5 15 2 US-09-069-827A-132 38 29 59.5 15 2 US-09-069-827A-132 39 21 56.8 6 1 US-08-459-568-20 40 21 56.8 6 1 US-08-399-411-20 41 21 56.8 6 2 US-09-586-472-20 42 21 56.8 6 2 US-09-586-472-20 43 21 56.8 8 1 US-08-459-568-75 44 21 56.8 8 1 US-08-459-568-75 45 21 56.8 8 2 US-09-516-859A-75 46 21 56.8 8 2 US-09-516-859A-75 47 21 56.8 8 2 US-09-516-859A-75 48 21 56.8 8 2 US-08-516-859A-75 49 21 56.8 8 2 US-09-516-859A-75 40 21 56.8 8 2 US-09-586-472-20 41 21 56.8 8 2 US-09-586-472-20 42 21 56.8 8 2 US-09-586-472-20 43 21 56.8 8 2 US-09-586-472-20 44 21 56.8 8 2 US-09-586-75 45 21 56.8 8 2 US-08-516-859A-75 46 21 56.8 8 2 US-08-516-859A-75 47 21 56.8 8 2 US-08-516-859A-75 48 21 56.8 8 2 US-08-516-859A-75
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Query Match

97.3%;

Score 36;

DB 2;

Length

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VLXDDLLEA 9

Best Local Matches

Similarity 8; Conserv

Conservative

88.9%;

Pred. No. 4.6e+05; 0; Mismatches 1;

Indels

0

Gaps

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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION UNMEER: US/09/489,760
CURRENT FILLING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILLNG DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 9
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US-09-269-250E-29
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US-09-489-760-1
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; NAME/KEY: PEPTIDE
; LOCATION: (3)
; LOCATION: (3)
; OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE
; OTHER INFORMATION: RESIDUE
US-09-489-760-1
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GENERAL INFORMATION:

APPLICANT: GOUING, Elsa

TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1

FILE REFERENCE: 58994

CURRENT APPLICATION NUMBER: US/09/269,250E

CURRENT FILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 38

SOFTWARE: PATENTIN Version 3.1

SEQ ID NO 29

LENGTH: 9

LENGTH: 9
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Best Local
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LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine (H) or an arginine
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ORGANISM: Human
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CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 9
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US-09-489-760-5
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RESULT 7
US-09-269-250E-26
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Best Local Similarity
Matches 8; Conserv
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Best Local Similarity
Matches 8; Conserv
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LENGTH: 9
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APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
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                                                                                                                                                                                                                               TYPE: PRT ORGANISM: KIAA0223 partial complementary DNA
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
0; Mismatches 1
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Sequence 26, Application US/09269250E Patent No. 6830883
GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa

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CURRENT APPLICATION NUMBER: US/09/269,250B
CURRENT FILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 28
LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
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US-09-269-250E-28
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APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION DATE: 1998-07-23
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Best Local S
Matches 8
                                                         NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver.
SEQ ID NO 14
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Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
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CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 26
                                                                                                                                                                                                                                                                                                               sequence 14, Application US/09489760 Patent No. 6878375
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Best Local Similarity
Matches 8; Conserv
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LENGTH: 13
TYPE: PRT
ORGANISM: Unknown Organism
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TYPE: PRT
ORGANISM: Artificial Sequence
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RESULT 11
US-09-269-250E-38
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APPLICANT: Hunt, Donald F

APPLICANT: Hard, Victor H

TITLE OF INVENTION: The HA-1 Antigen

FILE REFERENCE: 2183-4285US

CURRENT APPLICATION NUMBER: US/09/489,760

CURRENT FILING DATE: 2000-01-21

EARLIER APPLICATION NUMBER: PCT/NL98/00424

EARLIER FILING DATE: 1998-07-23

NUMBER OF SEQ ID NOS: 16
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US-09-489-760-16
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                                                                                                                                                                                   APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
                                                                                                                                                                                                                                                                                                                 Sequence 38, Applica
Patent No. 6830883
GENERAL INFORMATION:
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SEQ ID NO 16
LENGTH: 13
                                                              Matches
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Matches 8; Conserv
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APPLICANT: Rijksuniversiteit Te Leidin
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                                                                                                                                        TYPE: PRT
ORGANISM: HUMAN
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hes 8; Conservative
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                                                           $; Score 23; DB :
$; Pred. No. 4.6;
0; Mismatches
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Pred. No. 0.23;
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Pred. No. 0.23;
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                                                                           DB 2; L
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                                                                                         Length 8;
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                                                              Indels
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RESULT 12
US-08-940-095-104
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; MOLECULE TYPE:
US-08-940-095-104
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GENERAL INFOR
APPLICANT:
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Best Local Similarity
                                                                                                                                      Patent No. 6037323
GENERAL INFORMATION:
                                                                                                                                                        Patent No. 603732
                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                               APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: COTUZZÍ, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/940,095
FILING DATE: 29-SEP-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dufourcq, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
               APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOE
TITLE OF INVENTION: AND THEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: New York STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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1155 Avenue of the Americas
                                                                                                                                                                      Application US/08940093
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      650-493-5556
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Cornut, Isabelle
                                                                                                    Dasseux, Jean-Louis
Sekul, Renate
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                 APOLIPOPROTEIN A-I AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
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Pred. No. 1.4e+02;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/940,093
FILING DATE: 29-SEP-1997
CLASSIFICATION: 435
FRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 650-493-4935
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                                                                                                      COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: No. 6037323e
                                  SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                    TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 258
                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1155 AV CITY: New York STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM CO
OPERATING SYSTEM:
SOFTWARE: FastSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
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TELEFAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
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STREET: 1155 Avenue of the Americas
                                                                     COMPUTER: IBM CONTROL OPERATING SYSTEM:
                                                                                                                                                                               STATE: NY
                                                                                                                                                                                              CITY: New York
                                                                                                                                                                                                                STREET:
APPLICATION NUMBER: US/0 FILING DATE: 29-SEP-1997
                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                              104,
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                                                                                                                                                                                                                                                                                                                                                                                                                               6046166
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                                                                                                                                                                                                              B: Pennie & Edmonds LLP
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                     Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
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                    US/08/940,096
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Pred. No. 1.4e+02;
2; Mismatches 1
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CLASSIFICATION: 530 PRIOR APPLICATION NUMBER:

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US-09-465-719-104
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,093
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELECOMMUNICATION INFORMATION:
TELEPAX: 650-493-4935
TELEPHONE: 650-493-4935
TELEPAX: 650-493-5556
TELEEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
CTA NUTCENIEGO
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Sekul, Renate
APPLICANT: Ocrnut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
NUMBER OF SEQUENCES: 258
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PILING DATE:
ATTORNBY/AGENT INFORMATION:
NAME: COCUZZI, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0005-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEPAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: NY
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PILING DATE:
CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
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62.5%; Pred. No. 1.4e+02;
ative 2; Mismatches 1
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: No. 6265377e
US-09-465-719-104

Query Match
Best Local Similarity 62.5%; Pred. No. 1.4e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps
Qy 2 [INDLLEA 9
| ::|| | |
Db 2 LINDLLEA 9
Search completed: April 6, 2006, 17:05:27
Job time: 27.6667 secs
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Title:
Perfect score:
Sequence:
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No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     seq length: 0 seq length: 15
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Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Gapop 10.0 ,
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   Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
     GenCore version (c) 1993 - 2006
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US-11-007-740-20
US-11-007-740-20
US-11-007-740-24
US-10-623-176-43
US-10-623-176-65
US-10-623-176-74
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US-10-623-176-74
US-10-791-217-14
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RESULT 2 US-10-623-176-2

Sequence 2, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.

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Query Match Best Local Matches	TYPE: PRI ORGANISM: FEATURE: OTHER INF OTHER INF FEATURE: NAME/KEY: LOCATION: 10-623-176		RESULT 1 Sequence 1, Application US/1062317; Sequence 1, Application US/1062317; Publication No. US20040092446A1; Publicantion No. US20040092446A1; APPLICANT: Goulmy, Els A.J.M., APPLICANT: Hunt, Donald F., APPLICANT: Engelhard, Victor H.; APPLICANT ENGELBARG, Victor H.; TITLE OF INVENTION: HA-1 epitopes; FILE REFERENCE: 2183-6047US	30000044444000000000000000000000000000	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
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n Similarity 9; Conservat 1 VLXDDLLEA 9 1 VLXDDLLEA 9	Ö	T APPLICATION NUMBER: US/10/623,17 TF FILING DATE: 2003-07-18 APPLICATION NUMBER: 09/489,760 FILING DATE: 2000-01-21 APPLICATION NUMBER: EP 97202303.0 FILING DATE: 1997-07-23 APPLICATION NUMBER: PCT/NL98/00424 APPLICATION NUMBER: JP 2000-504165 FILING DATE: 1998-07-23 APPLICATION NUMBER: JP 2000-504165 FILING DATE: 2000-01-24 OF SEQ ID NOS: 77 RE: PatentIn Ver. 2.1 NO 1 H: 9	ion 040 Els Els HA,	ស់សំធំធំតំតត់តំ	
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APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof

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US-10-623-176-2
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
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Publication No. US20040092446A1
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                          FEATURE:
NAME/KEY: SITE
LOCATION: (1)..(9)
                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
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Pred. No.
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                                                      Score 36; DB 4;
Pred. No. 1.7e+06;
                                      Mismatches
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US-10-791-217-2
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILLING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILLING DATE: 2000-01-21
PRIOR FILLING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
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Publication No. US20040191268A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2
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Best Local Similarity 88.9%; Pred. No. 1.7e+06; Matches 8; Conservative 0: Migmatches
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
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NAME/KEY: MISC_FEATURE
LOCATION: (3)..(3)
OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
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                                                                                                                                                                                                                                                                                 ORGANISM: Artificial sequence FEATURE:
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US-10-791-217-5
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APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                    Query Match
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Matches 8
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APPLICANT: GOULDY, Elsa A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot
TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
CURRENT APPLICATION NUMBER: US/10/861,335
CURRENT FILLING DATE: 2004-06-04
CURRENT FILLING DATE: 2004-06-04
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                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1 SEQ ID NO 1
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LENGTH: 9
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                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial sequence
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                                  VLXDDLLEA 9
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                                                                                     97.3%;
88.9%;
                                                                Pred. No. 1.7e-
0; Mismatches
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 4; 1
                                                                                                    DB 5;
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                                                                                                    Length 9;
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                                                                  <u>.</u>
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                                                                  Gaps
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                                                                  0
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RESULT 10
US-11-007-740-29
; Sequence 29, Application US/11007740
; Publication No. US20050233350A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-11-007-740-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-11-007-740-20
; Sequence 20, Application US/11007740; Publication No. US20050233350A1; GENERAL INFORMATION:
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US-11-007-740-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 8
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 1094-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
CORTWANDE DESCRIPTION OF THE CONTROL O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.1 SEQ ID NO 20
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Goulmy, E18a
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 8; Conserv
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Pred. No. 1.7e
0; Mismatches
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Pred. No. 1.7e+06;
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APPLICANT: GOLINY, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR APPLICATION NUMBER: JP 97202303.0
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 1908-07-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 24
LENGTH: 10
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US-10-623-176-24
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                                                                                                            RESULT 12
US-10-623-176-43
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PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOPTWARE: PatentIn version 3.1
SEQ ID NO 29
Sequence 43, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 97.3%; Score 36; DB Best Local Similarity 100.0%; Pred. No. 1. Matches 9; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                                           Matches
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TYPE: PRINTER:
ORGANISM: Human
FEATURE:
FEATURE:
FEATURE:
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine (H) or an arginine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide OTHER INFORMATION: wherein X can be H or R
                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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hes 0;
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APPLICANT: Goulmy, B1s A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLS OF INVENTION: HA-1 epitopes and uses thereof
FILE REPERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER: DF 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER: DF 2000-504165
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                                                                                                                               ; NAME/KEY: SITE
; LOCATION: (1)..(10)
US-10-623-176-44
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US-10-623-176-44
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; NAME/KEY: SITE
; LOCATION: (1)..(10)
US-10-623-176-43
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PC7/NL98/00424
PRIOR APPLICATION NUMBER: DF 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
                                                                                                                                                                                                                                                                                      SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 44
LENGTH: 10
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 44, Application US/10623176 Publication No. US20040092446A1
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                                             Matches
                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                               FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE:
                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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    1 VLXDDLLEA 9
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8; Conservative
                                             Conservative
                                                               97.3%;
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                                                                      Score 36; DB Pred. No. 1.8;
                                               Mismatches
                                                                                         DB 4;
                                                                                         Length 10;
                                                  Indels
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                                                  Gaps
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RESULT 14 US-10-623-176-65 皮

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PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 65
                                                                                                                                                                                                                                                                                                                                                                               RESULT 15
US-10-623-176-74
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Publication No. US20040092446A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 65, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE: NAME/KBY: SITE LOCATION: (1)..(12)
:-10-623-176-65
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
TENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 VLXDDLLEA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                           VLRDDLLEA 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 36;
Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
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Search completed: April Job time: 97.6667 secs
                                                                                                                                                                       US-10-623-176-74
                                                                                                                             Query Match
Best Local Similarity
                                                                                                                 Matches
                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: KIAA0223 OTHER INFORMATION: sequence derived from a presumed HA-1 negative OTHER INFORMATION: individual FEATURE:
                                                                                                                                                                                 NAME/KEY: SITE LOCATION: (1)..(13)
                                                                       1 VLXDDLLEA 9
                                                                                                               8; Conservative
                                                        3 VIRDDLLEA 11
             6, 2006, 17:28:45
                                                                                                                              97.3%;
88.9%;
                                                                                                                           Score 36; 1
Pred. No. 2
                                                                                                                 Mismatches
                                                                                                                                         DB 4;
                                                                                                              1; Indels
                                                                                                                                         Length 13;
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                                                                                                               Gaps
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Result
No.
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Maximum
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
BG
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              Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/USO1_NEW_PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/USO1_NEW_PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/USO1_NEW_PUB.pep:*
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Gapop 10.0 , Gapext 0.5
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37
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                                   VLXDDLLEA 9
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Copyright (c) 1993 - 2006 Biocceleration
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                                                        DB
                                                                      US-11-010-748A-11
US-11-010-748A-12
US-11-12-740-1404
US-10-927-435-63
US-10-927-634-63
US-10-927-634-63
US-11-129-143-169
US-11-129-143-171
US-11-129-143-171
US-11-129-143-171
US-11-129-143-173
US-11-129-143-173
US-11-129-143-173
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US-11-129-143-173
US-11-129-143-173
US-11-129-143-173
US-11-129-143-173
US-11-09-143-173
US-11-09-173-159
US-11-09-173-159
US-11-09-173-173
US-11-09-173-173
                              US-11-225-686-2954
US-11-225-686-3877
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Sequence 11, Appl Sequence 120, Appl Sequence 130, Appl Sequence 71, Appl Sequence 170, Appl Sequence 170, Appl Sequence 171, Appl Sequence 171, Appl Sequence 172, Appl Sequence 173, Appl Sequence 173, Appl Sequence 173, Appl Sequence 65, Appl Sequence 65, Appl Sequence 189, Appl Sequence 190, Appl Sequence 102, Appl Sequence 102, Appl Sequence 102, Appl Sequence 103, Appl Sequence 103, Appl Sequence 103, Appl Sequence 2954, Appl Sequence 2954, Appl Sequence 3974, Appl Sequence 397
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RESULT 2 US-11-010-748A-12

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Sequence 12, Application US/11010748A
Publication No. US20050244421A1
GENERAL INFORMATION:
APPLICANT: MOCK Patent GmbH
APPLICANT: STRITTWATTER, Wolfgang
APPLICANT: MOLL, Heidrun
APPLICANT: SCHARM, Burkhard

Oy Mai	SEQ 11. SEQ 11	SP S	BR CITI.	
Matches 1	ID N ID N ID N INGTH IPE: IPE: IPE: IPE: IPE: IPE: IPE: IPE:	-11-010-748 Sequence 11 Publication GENERAL INF APPLICANT: APPLICANT: APPLICANT: ITIFLE OF 11 FILE REFERI CURRENT FILE PRIOR APPLI PRIOR APPLI PRIOR FILLI		44444000000000000000000000000000000000
1 1 85	WE PATENTIN OUT 1 H: 9 PRT ISM: Homo Bap EX: MISC FEA CGY: MISC FEA CON: (1)9) INFORMATION: INFORMATION: 1.1FORMATION: 748A-11	11, 11, 11, 11, INFOR NT: WIT: WIT: NT: WIT: FILI PPLIC PPLI		1199
Similarity 8; Conser VLXDDLLEA VLHDDLLEA	entin vers no sapiens SC FEATURE 1 (9) ATION: Hum	11-010-748A-11 11-010-748A-11 11-010-748A-11 UNDICATION NO. US20050244 ENERAL INFORMATION: ENPELCANT: METCH PATENT APPLICANT: STRITTMATTER, APPLICANT: MOLL, Heidrun APPLICANT: SCHARM, Burkh TITLE OF INVENTION: COMPO FILE REFERENCE: MER-136 CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2003-0 PRIOR APPLICATION UNMBER: PRIOR FILING DATE: 2003-0		551114444444444444444444444444444444444
88. vative	Homo sapiens Homo sapiens MISC FEATURE (1) - (9) ORMATION: Human of the community of the	R G A A C C C C C C C C C C C C C C C C C		113311331131131131131131131131131131131
" 'o 'i	on 3.1 n minor epitopes			011110000111111111111
Pred. No. 1.4e+05 0; Mismatches	histocompa	Application US/11010748A Application US/11010748A Io. US20050244421A1 MATION: MATION: MATION: MATION: MATION: MOLL, Heidrun SCHARM, Burkhard ENTION: COMPOSITIONS AND METHODS CE: MER-136 CE: MER-136 ICATION NUMBER: US/11/010,748A NG DATE: 2004-12-13 ATION NUMBER: PCT/EP03/06251 DATE: 2003-06-13 ATION NUMBER: EP02013423.5 DATE: 2002-06-13 Q ID NOS: 926	ALIGNMENTS	US-11-225-686-3915 US-11-225-686-3932 US-11-225-686-3941 US-11-202-009-2954 US-11-202-009-3915 US-11-202-009-3916 US-11-202-009-3916 US-11-202-009-3916 US-11-202-009-3931 US-11-202-009-3931 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-203-137-7 US-11-203-137-7 US-11-203-137-7 US-11-203-139-1739 US-11-207-078-93 US-11-207-078-93 US-11-207-078-93
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Сарв	characterized	RESPONSE		115, 312, 314, 315, 316, 316, 317, 317, 317, 317, 317, 317, 317, 317
0;	rized	88		

FEATURE: misc_feature OTHER INFORMATION: Utility:

Useful

for

making

plants

sterile

and

for

genetic

confinen

flowers

for making taller plants

and plants with longer

NAME/KEY: misc_feature

OTHER INFORMATION: Utility: Useful for making ornamental plants with modified

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US-11-010-748A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-11-172-740-1404
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR TITLE OF INVENTION: PLANT CHARACTERISTICS AND PHENOTYPES FILE REFERENCE: 2750-1602PUS2 FILE REFERENCE: 2750-1602PUS2 FILE REFERENCE: 2750-1602PUS2 FILE REPLICATION NUMBER: US/11/172,740 CURRENT FILING DATE: 2005-06-30 PRIOR APPLICATION NUMBER: 60/583,621 PRIOR APPLICATION NUMBER: 60/583,621 PRIOR FILING DATE: 2004-06-30 PRIOR FILING DATE: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2004-06-30 PRIOR APPLICATION NUMBER: 60/584,800 PRIOR FILING DATE: 2004-06-30
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/584,829
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 2523
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FEATURE:
NAME/KEY: misc feature
...(258)...(258)
...(258)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Arabidopsis thaliana
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (208)..(258)
OTHER INFORMATION: Pfam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (208)..(258)
OTHER INFORMATION: Pfam Name:
                                                                                OTHER INFORMATION: Utility: Useful for making larger plants
                                                                                                                                                                                                                                    OTHER INFORMATION: Utility:
                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature LOCATION: (208)..(258)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                    NAME/KEY: misc_feature
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8; Conserva
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88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                             Name:
                                                                                                                                                                                                                                    Useful
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Pred. No. 1.4e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                 PHD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam
                                                                                                                                                                                                                                                                                                                                                                                                 Pfam
                                                                                                                                                                                                                                    for increasing chlorophyll and photosynthetic cap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Description: PHD-finger
                                                                                                                                                                                                                                                                                                                                                                                                 Description: PHD-finger
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US-10-927-435-63
; Sequence 63, Application US/10927435
; Publication No. US20060045884A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                RESULT 5
                                                                                                 APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: ANTI-CANCER VACCINES
FILE REFERENCE: UTSC: 788USC1
CURRENT APPLICATION NUMBER: US/10/927,634
CURRENT FILING DATE: 2004-08-26
PRIOR APPLICATION NUMBER: 60/489,238
PRIOR FILING DATE: 2003-08-26
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       Sequence 63, Application US/10927634 Publication No. US20060045881A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                        SEQ ID NO 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: VACCINES FOR AUTOIMMUNE AND INFECTIOUS DISEASE
FILE REFERENCE: UTSC: 788USC2
CURRENT APPLICATION NUMBER: US/10/927,435
CURRENT FILING DATE: 2004-08-26
PRIOR APPLICATION NUMBER: 60/489,238
PRIOR APPLICATION NUMBER: 60/489,238
PRIOR PLING DATE: 2003-08-26
PRIOR PLING DATE: 2003-08-26
PRIOR PLING DATE: 2003-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Utility: Useful OTHER INFORMATION: inflorescences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BNGTH:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                     ORGANISM: Artificial Sequence
                                                      TYPE: PRT
                       FEATURE:
                                                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Description of Artificial Sequence: Peptide
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Pred. No. 1.3e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 21;
Pred. No.
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US-10-353-783-71
US-10-353-783-71
; Sequence 71, Application US/10353783
; Publication No. US20050261175A1
; Publication No. Wrisztina M
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 71:
US-10-353-783-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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Matches 5; Conserv
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                                                                                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION UNMBER: U8/10/353,783
PRICATION UNMBER: U8/10/353,783
PRICATION UNMBER: U8/10/353,783
PRIOR APPLICATION UNMBER: U8/10/353,783
PRIOR APPLICATION UNMBER: 08/448,729
PRIOR APPLICATION UNMBER: 08/448,729
PILING DATE: 24-MAY-1995
APPLICATION UNMBER: 08/172,329
PILING DATE: 21-DEC-1993
APPLICATION NUMBER: 07/982,255
PILING DATE: 25-NOV-1992
PILING DATE: 25-NOV-1992
PILING DATE: 25-NOV-1992
PILING DATE: 25-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
TELEFAX: 312/474-0448
                                                                                                                                                                                                                                                                                                                                             FILING DATE: 11-JUN-1990
APPLICATION NUMBER: 07/422,383
FILING DATE: 16-OCT-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole,
STREET: 6300 Sears Tower, 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suggs, Sidney V.

Martin, Francis H.

TITLE OF INVENTION: Stem Cell Factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Zsebo, Krisztina M.
Bosselman, Robert A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 VLXDDLLE 8
                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 01017/32958A
                                                                                                                                                                                                                                                                                                  NAME: Clough, David W. REGISTRATION NUMBER: 36,107
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/589,701
FILING DATE: 01-OCT-1990
APPLICATION NUMBER: 07/573,616
FILING DATE: 24-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: United States of America ZIP: 60606-6402
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STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/537,198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/684,535 FILING DATE: 10-APR-1991
                                                                                                                              LENGTH: 15 amino acids
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Pred. No.
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South Wacker Drive
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APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: HUMBELIN, Markus
APPLICANT: HOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: WELISEEV, Alexed A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOPTWARE: Patentin version 3.1
SEQ ID NO 168
                                                                                                                                  US-11-129-143-169
                                                                                                                                                                                                                                                                                                                                                                                                                                   US-11-129-143-169
; Sequence 169, Application US/11
; Publication No. US20050266518A1
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US-11-129-143-168
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                                                                  Matches
                                                                                Query Match
Best Local :
                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 50.0
Matches ... 4; Conservative
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                                                                                                                                                                                                                                 FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                 APPLICANT: LOPEZ-ULÍBARRI, RUAL
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, J
                                                                                                                                               TYPE: PRT
ORGANISM: Rhizobium
                                                                                                                                                                                   LENGTH: 5
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les 3; Conserv
1 DDILD 5
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                                 4 DDLLE 8
                                                                                Similarity
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                                                                  Conservative
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5. US20050266518A1
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                                                                                                                                               sp. strain NGR234
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                                                                              54.1%;
                                                              Score 20; DB 7; Le
Pred. No. 1.4e+05;
2; Mismatches 0;
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Pred. No. 1.4e+05;
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Pred. No. 2.1e+02;
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                                                                                            Length 5;
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APPLICANT: BRETŽEL, Werner
APPLICANT: HUMBELIN, MARKUS
APPLICANT: HUMBELIN, MARKUS
APPLICANT: MAYER, Anne F.
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION NUMBER: US/11/129,143
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 170
                       US-11-129-143-172
; Sequence 172, Application US/11129143
; Publication No. US20050266518A1
; GENERAL INFORMATION:
; APPLICANT: BERRY, Alan
nnownyor. Warney
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BERRY, Alan
APPLICANT: BERTZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: WAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
APPLICANT: YELISEEV, Alexei A.
APPLICANT: CISEVS Alexei A.
APPLICANTON: IMPROVED ISOPRENOID PRODUCTION
FILLE REFERENCE: C18435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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; TYPE: PRT
; ORGANISM: Bacillus stearothermophilus
US-11-129-143-170
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                                                                                                                          RESULT 11
                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Bacillus subtilis US-11-129-143-171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 170, Applic Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                 SOPTWARE: PatentIn version 3.1
SEQ ID NO 171
LENGTH: 5
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Best Local :
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                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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3; Conserva
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3; Conserv
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HUMBELIN, Markus
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10. US20050266518A1
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Pred. No. 1.4e+05;
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Pred. No.
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APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 172
LENGTH: 5
TYPE: PRT
ORGANISM: Escherichia coli
US-11-129-143-172
                                                        Sequence 126, Application US/10982440

Publication No. US20060018909A1

GENERAL INFORMATION:

APPLICANT: Oliner, John
APPLICANT: Graham, Kevin
ITILE OF INVENTION: Angiopoietin-2 Specific Binding Agents
FILE REFERENCE: 04-881-A

CURRENT APPLICATION NUMBER: US/10/982,440

CURRENT FILING DATE: 2004-11-04

PRIOR APPLICATION NUMBER: 60/620,161

PRIOR FILING DATE: 2004-10-19

JENIOR FILING DATE: 2004-10-19
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US-10-982-440-126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 173, Application US/11129143 Publication No. US20050266518A1
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APPLICANT:
SOFTWARE: PatentIn version 3.3 SEQ ID NO 126 LENGTH: 7
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Best Local S
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APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REPERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Haemophilus influenzae
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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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RESULT 15

US-11-021-305-65.

US-11-021-305-65.

Sequence 65, Application US/11021305

Publication No. US20050282733A1

GENERAL INFORMATION:

APPLICANT: Prins, Johannes B

APPLICANT: Hutley, Louise J

TITLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI169.001CP1

CURRENT FILING DATE: US/11/021,305

CURRENT FILING DATE: 2004-12-23

PRIOR APPLICATION NUMBER: US/11/021,305

CURRENT FILING DATE: 2002-06-27

NUMBER OF SEQ ID NOS: 170

SOFTWARE: PATENTIN VERSION 3.3

SEQ ID NO 65

LENGTH: 8

TYPE: PRT

ORGANISM: mammalian

US-11-021-305-65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Prins, Johannes B
APPLICANT: Hutley, Louise J
FITTLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI69.001CP1
CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILLNG DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR PILLING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.3
SEQ ID NO 62
TYPE: PRT
CORGANISM: mammalian
US-11-021-305-62
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; ORGANISM: Homo sapiens
US-10-982-440-126
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US-11-021-305-62
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                                                                        Query Match 54.1%; Score 20; DB 7; Les Best Local Similarity 100.0%; Pred. No. 1.4e+05; Matches 4; Conservative 0; Mismatches 0;
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Result No.	Score	% Query Match	Length	DB	ID	Description
_	45	100.0	9	2	٠	Aaw99196
2	45	100.0	9	N	AAW97375	Aaw97375
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7	37	82.2	9	N	AAW97374	Aaw97374
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11	36	80.0	9	N	AAW99195	Aaw99195
12	36	0	9	ນ	AAW97572	Aaw97572
13	31	8	9	œ	ADU99853	Adu99853
14	30	66.7	9	N	AAW97373	Aaw97373
15	29	64.4	13	4	AAU02706	Aau02706
16	28	62.2	9	N	AAY10122	Aay10122
17	28		9	σ	ABG79805	Abg79805
18	28	62.2	9	8	ADK68732	Adk68732
19	28	•	9	œ	ADK05291	Adk05291
20	28	•	9	œ	ADK05293	Adk05293
21	28	'n	9	œ	ADQ10530	Adq10530
22	28	ە:	9	8	ADS81010	Ads81010
23	28	62.2	10	8	ADK05295	Adk05295
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Sequence 9 diseases

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57.8	57.8	57.8	57.8			57.8	57.8	57.8	57.8	57.8	57.8	57.8	57.8	57.8	60.0	•	60.0	60.0	60.0	62.2
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-	Abp71119 E1	Aaw73901 B.	Adu99636 BE	Adt49074 Ht	Adt73023 Hu	Adm96258 Hu	Adm96261 Hu	Aae31275 Hu		Ady63770 Ht	Abu97231 Er	Abu97234 Er	Abu97232 Er	Abu97233 Er	Aab85066 Ht		Adq26722 To	Adt02787 Ht	Abj20115 M	Add23269 B1
HLA bindi	E10 prote	fo	BFA5 tumo	Human BFA	Human RSV	Human ser	Human ser	Human mag		Human alb			Enzyme pe	Enzyme pe	Human ser	Human ser	Topo V Hh	Human tum	MHC bindi	Breast ca

ALIGNMENTS

AAW99196;

AAW99196 standard; peptide; 9 AA.

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RESULT 1
AAW99196
ID AAW9
XX AAW9
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PT disk
                                                                                                                  The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 32; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-153312/13.
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DB 2;

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R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant;
severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-i-dictoria a colla anti-i-dictoria a co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. det of genetic aberrances.
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02-JUN-1998;
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ADH,40333;
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                                                                                   ADH40333 standard; peptide; 9 AA
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9; Conserve
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98EP-00870125.
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Pred. No. 2e+06;
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Pred. No. 2e+06;
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RESULT 4
AAW99199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-2002; 2002EP-00013423
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                                                                                                                                                AAW99199;
                                                                                                                                                                                    AAW99199 standard; peptide;
                                                                          vR cell KIAA0223 protein sequence.
                                                                                                             20-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                Similarity
9; Conserve
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100.0%; Pred. No. 2e-
tive 0; Mismatches
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2e+06;
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RESULT 5
AAW97415
BX5XXFXBXBX8XgXgXGXFXGXGXGX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases. The present sequence represents a KIAA0223 sequence given in the present invention
         Goulmy EAJM,
                                                                                                      04-FEB-1999.
                                                                                                                            W09995173-A1
                                                                                                                                                                         T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease; neoplastic haematopoietic cell; KIAA0223 polymorphism.
                                                                                                                                                                                                                           KIAA0223
                                                                                                                                                                                                                                                   20-MAY-1999
                                                                                                                                                                                                                                                                                                AAW97415 standard; protein; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
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                                                                                                                                                  Homo sapiens.
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                                 (UYLE-) RIJKSUNIV LEIDEN.
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         Hunt DF,
                                                        97EP-00202303
                                                                              98WO-NL000424.
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                                                                                                                                                                                                                          in HA-1
        Engelhard VH;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Score 45;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        versus graft disease in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HA-1 - useful to treat immune
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Best Local Similarity
The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                              A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents the KIAA0223 polymorphism in HA-1 positive homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
                                                                                                                     Disclosure; Page 15;
                                                                                                                                                                                                                   WPI; 1999-153312/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Minor histocompatibility antigen HA-1 T-cell epitope #3
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N-PSDB; AAX16081.
                                                                                                                                                                                                                                                    Goulmy EAJM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              versus-host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
                                                                                                                                                                                                                                                                                      (UYLE-) RIJKSUNIV LEIDEN
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                                                                                                                                                                                                                                                    DF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11
                                                                                                                   47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 45;
100.0%; Pred. No.
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                                                                                                                                                                 HA-1 - useful to treat i versus graft disease in
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Best Local
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                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases
                                                                                                                     The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-MAY-1999
                                                                                                                                                                                                                                                                                 Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. det of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                             23-JUL-1997;
02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-FEB-1999
                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                      WPI; 1999-142960/12
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                                                            Local Similarity
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8; Conserv
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                       VLHDDLLEA 9
VLRDDLLEA
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98EP-00870125.
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Pred. No.
                                                            Score 37; DB 2;
Pred. No. 2e+06;
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                                                 Mismatches
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2e+06;
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ADH403A
ADH403A
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RESULT 9
AAW99198
ID AAW9
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AC AAW9
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AC AAW9
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AC AAW9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method for providing epitopes of allelic variants of antigenic proteins from specific species based on single nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein, identifying, selecting allelic peptide/protein variants, expression product or its fragment encoded by DNA sequence having SNP, creating variant epitopes, selecting epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is useful for generating a SNP profile of one or more individuals from a given species by applying the method for several protein from the concer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease. A method of the invention can be progression, regression or onset of a treated disease. The present contains and treating cancer, preferably leukaemia, and for determining the progression, regression or onset of a treated disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-082200/08.
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sequence is used in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9 AA;
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 37;
Pred. No.
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2e+06;
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20-MAY-1999

(first entry)

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AAW99198 standard; peptide; 13

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Best Local
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                                                                                                                               T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis neoplastic haematopoietic cell; KIAA0223 polymorphism.
                                                                                                                                                                                                              20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 31; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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 23-JUL-1997;
                          23-JUL-1998;
                                                                              WO9905173-A1
                                                                                                      Homo sapiens
                                                                                                                                                                                   KIAA0223 polymorphism in HA-1 negative homozygous individuals.
                                                                                                                                                                                                                                                               AAW97414 standard;
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8; Conserv
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97EP-00202303
                           98WO-NL000424
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                   5.1;
                                                                                                                                           Disease; autoimmune disease;
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RESULT 11
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Best Local S
Matches 8
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A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone
                                                                                                                                                                                                                                                                                                                        Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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N-PSDB; AAX16080.
                                                                 Goulmy EAJM,
                                                                                                                                                                            04-FEB-1999
                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                              Minor histocompatibility antigen HA-1 T-cell epitope #1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW99195 standard;
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                                                                                                                       23-JUL-1997;
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                                                                                                                                                                                                                                             Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
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                                                                                            (UYLE-) RIJKSUNIV LEIDEN
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les 8; Conserv
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                                                                 Hunt DF,
                                                                                                                       97EP-00202303
                                                                                                                                                 98WO-NL000425
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marrow and organ transplantation

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease; neoplastic haematopoietic cell.
                                              The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides.
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Misc-difference
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                                                                                                                                                                                                                                                               Claim 1;
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                                                                                                                                                                                                                                                                                                                                        Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention comprises an expression vector that contains a nucleic acid encoding a tumor antigen (e.g. BCZ4 or BPY3). The expression vector of the invention is useful for the expression of multiple tumor antigens and the prevention or treatment of cancer. The present amino acid sequence represents a BCZ4 tumor antigen antigenic peptide that was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New expression vector comprising a nucleic acid encoding a tumor antigen, e.g. BFA4, BCY1, BFA5, BCZ4, or BFY3, useful for expressing multiple tumor antigens, or for preventing or treating cancer.
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Singh-Sandhu D;
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16-MAY-2003; 2003US-0471193P.
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                                                                                                                                                                                                                                                                                             The present sequence represents an epitope of the minor histocompatibility antiqen HA-1. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antiqen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and HA-1 peptides can be used anti-idiotypic B cells and/or T cells and
Antibody; adipocyte; heavy chain; light chain; obesity; fat;
                       CDR region of anti-adipocyte antibody heavy chain, FAT 71.
                                                  29-AUG-2001
                                                                           AAU02706;
                                                                                              AAU02706 standard; peptide;
                                                                                                                                                                                                                                                               Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Typing minor histocompatibility antigen HA-1 - identifying HA-1 R and HA-1 H alleles of HA-1,
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CC AAU02501-AAU02635, and AAU02641-AAU02748 represent the amino acid CC sequences of anti-adipocyte monoclonal antibody heavy chain, light chain, CC and heavy chain complementarity determining regions (CDR) of the CC invention. The antibodies can be used in the treatment of obesity and CC obesity related diseases. The antibodies can be used to deliver drugs or CC pro-drugs directly to the fat mass of an obese patient or the antibody CC can be used as a therapeutic itself. Antibodies binding specifically to CC adipocytes can be used to activate the immune system to destroy the cells CC by complement mediated lysis. The antibodies may be labeled with a CC used in methods of diagnosis in human subjects e.g. to determine the presence or level of adipocytes in a cell or tissue sample. CC determine the presence or level of adipocytes in a cell or tissue sample. CC The antibodies can be used as an alternative means of treatment for obese patients other than undergoing surgery to remove excess fat. Antibodies CC abdominal fat associated with heart disease
   Query Match
Best Local Similarity
Matches 5; Conserv
                                                                               Sequence 13 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7; Page 78; 182pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diseases.
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Qy 2 LHDDLLEA 9 : Db 3 VHDVTLHA 10	Query Match 42.2%; Score 19; DB 2 Best Local Similarity 50.0%; Pred. No. 1.7e+ Matches 4; Conservative 1; Mismatches	RESULT 2 S30608 translation elongation factor eEF-2 - bovine (fragment) C;Species: Bos primigenius taurus (cattle) C;Accession: \$30608 R;Srown, B.A.; Bodley, J.W. FRBS Lett. 103, 253-255, 1979 A;Title: Primary structure at the site in beef and wheat A;Reference number: \$30607; MUID:79236796; PMID:467667 A;Accession: \$30608 A;Molecule type: protein A;Residues: 1-15 <bro> A;Cross-references: UNIPARC:UPI000017482A A;Experimental source: liver C;Keywords: GTP binding; protein biosynthesis</bro>	Qy 1 VIHDDL 6 :: Db 10 IIEDDL 15	Query Match 46.7%; Score 21; DB 2; Best Local Similarity 50.0%; Pred. No. 6.9e+02; Matches 3; Conservative 2; Mismatches 1	C; Date: 01-DeC-1995 #sequence revision 01-DeC-1995 #text_C; Accession: A53594 R; Lenter, M.; Vestweber, D. J. Biol. Chem. 269, 12263-12268, 1994 A; Title: The integrin chains beta-1 and alpha-6 associate A; Reference number: A53594; MUID:94216347; PMID:8163531 A; Accession: A53594 A; Status: preliminary A; Molecule type: protein A; Residues: 1-15 < LENY A; Residues: 1-15 < LENY A; Cross-references: UNIPROT:Q7M063; UNIPARC:UPI000017C62E C; Keywords: endoplasmic reticulum; molecular chaperone	ALI	30 16 35.6 14 2 S29789 31 16 35.6 15 2 S48685 32 16 35.6 15 2 B32800 33 16 35.6 15 2 B32800 34 16 35.6 15 2 B26501 35 15 33.3 7 2 X59489 37 15 33.3 11 2 X14454 38 15 33.3 12 PT0256 40 15 33.3 13 2 PT0256 41 15 33.3 13 2 S65612 41 15 33.3 14 2 X61306 42 15 33.3 14 2 PM0009 45 15 33.3 15 2 PA0009 45 15 33.3 15 2 PA0009
	DB 2; Length 15; .7e+03; ss 3; Indels 0; Gaps 0;	(fragment) -1997 #text_change 07-May-1999 f and wheat elongation factor 2 of ADP-ribo ID:467667		Length 15; 12; 1; Indels 0; Gaps 0;	2-1995 #text_change 09-Jul-2004 11-6 associate with the chaperone calnexin pr 11D:8163531 PPI000017C62E Chaperone	# T P P P P P P P P P P P P P P P P P P	hypothetical prote extension protein 3-dehydroquinase - hypothetical protein lipase ribosomal protein lipase ribosomal protein forpotein kinase C i 6-phosphofructokin hypothetical protein hypothetical protein thoulin alpha-chai photosystem I 13K Ig H chain V-D-J r ribonuclease M (EC seed storage prote self-incompatibili

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peptide-N4-(N-acetyl-beta-glucosaminyl)asparagine amidase (EC 3.5.1.52) A, N;Alternate names: peptide N-glycosidase C;Species: Prunus dulcis var. sativa (sweet almond) C;Date: 19-May-2000 #sequence_revision 19-May-2000 #text_change 09-Jul-2004 C;Accession: B59272
                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
B59272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R; Priedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984
A; Title: Characterization of genomic clones specifying rabbit A; Reference number: 146868; MUID:84221901; PMID:6328491
A; Accession: 146868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPARC:UPI0000170C44; GB:S51512; NID:g261982; A;Experimental source: GH3 anterior pituitary cells A;Note: sequence extracted from NCBI backbone (NCBIP:120927)
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J. Biol. Chem. 267, 25703-25708, 1992
A;Title: Two isoforms of the thyrotropin-releasing hormone receptor generated A;Reference number: A45096; MUID:93100278; PMID:1334485
A;Accession: A45996
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                                                                                                                                                                                    R;Altmann, F.; Paschinger, K.; Dalik, T.; Vorauer, K. Eur. J. Biochem. 252, 118-123, 1998
A;Title: Characterisation of peptide-N4-(N-acetyl-bet: A;Reference number: A59272; MUID:98181894; PMID:952377
A;Rccession: B59272
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    alpha-myosin heavy chain - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 09-Jul-2004
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A; Residues: 1-7 < FRI>
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50.0%;
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Pred. No.
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S29485
GTP-binding
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C26997
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                                                                                                                                                                                         PS I complex subunit 8 - cucumber (fragment)
C;Species: Cucumis sativus (cucumber)
C;Date: 05-Jan-1996 #sequence_revision 05-Jan-1996 #text_change 09-Jul-2004
C;Accession: E56819
R;Iwasaki, Y.; Ishikawa, H.; Hibino, T.; Takabe, T.
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                                                                                                                                 R;Iwasaki, Y.; Ishikawa, H.; Hibino, T.; Takabe, T. Biophim. Biophys. Acta 1059, 141-148, 1991
A;Title: Characterization of genes that encode subunits A;Reference number: A56819; MUID:91355209; PMID:1883835
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                                                   A; Molecule type: protein A; Residues: 1-15 < IWA>
                                                                                                                                                                                                                                                                                             E56819
                                                                                           A;Status: preliminary
                                                                                                          A;Accession: E56819
                                                                                                                               A; Reference number:
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                             UNIPROT:P42052; UNIPARC:UPI0000132593
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R;Graves, P.B.; Kaminsky, L.S.; Halpert, J. Biochemistry 26, 3887-3894, 1987 A;Title: Evidence for functional and structural multiplicity A;Reference number: A26997; MUID:88000604; PMID:3651420 A;Accession: C26997
                                                                                                                                                                                                                         FEBS Lett. 317, 53-56, 1993
A;Title: Association of three small GTP-binding proteins with A;Reference number: S29485; MUID:93154521; PMID:8428634
A;Accession: S29485
                                                                                                                                                                                                                                                                                                GTP-binding protein o-ral - Pacific electric ray (fragment) C;Species: Torpedo californica (Pacific electric ray) C;Date: 22-Nov-1993 #sequence_revision 27-Feb-1997 #text_cha C;Accession: S29485 C;Accession: S29485 R;Volknandt, W.; Pevsner, J.; Elferink, L.A.; Scheller, R.H. FEBS Lett. 317, 53-56, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPROT:Q06884; UNIPARC:UPI0000174DA8
C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase; oxidoreductase; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 31-Dec-2004
C;Accession: C26997
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C;Species: Rattus norvegicus (Norway rat)
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                                                                                                                                   A;Molecule type: protein
A;Residues: 1-15 <VOL>
A;Cross-references: UNIPARC:UPI000017BF33
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2 LHDDLLE 8
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Pred. No.
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                                                                                          Length 15;
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C;Accession: S55696
R;Hunt, M.; Koehler, P.
Biochim. Blophys. Acta 1249, 15-22, 1995
A;Title: Purification and characterization
A;Title: number: S55696; MUID:95284106;
virG protein - Agrobacterium sp. (fragment)
() Species: Agrobacterium sp.
() Species: Agrobacterium sp.
() Date: 19-Mar-197 #sequence_revision 01-Aug-1997 #text_change 09-Jul-2004
() Accession: $13224
R;Tamamoto, S.; Aoyama, T.; Takanami, M.; Oka, A.
J. Mol. Biol. 215, 537-547, 1990
A;Title: Binding of the regulatory protein VirG to the phased signal sequen
A;Reference number: $13224; MUID:91039316; PMID:2231718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 10
S55696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   phosphoenolpyruvate carboxykinase - Trypanosoma brucei
C;Species: Trypanosoma brucei
C;Date: 28-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 09-Jul-2004
C;Accession: S55696
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A;Molecule type: protein
A;Residues: 1-9 <HUN>
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C;Superfamily: cytochrome P450 homology
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2; Conserve
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Pred. No. 2.6e+03;
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Pred. No.
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                    signal sequences upstream
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cytochrome P450 UT-7b - rat (fragment)
(;Species: Rattus norvegicus (Norway rat)
C;Date: 19-May-1994 #sequence_revision 10
C;Accession: S39762
R;Ohishi, N.; Imaoka, S.; Suzuki, T.; Fun
                                                                                                     RESULT
S39762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eur. J. Blochem. 213, 591-597, 1993
A;Title: Characterization of a clotting protein, isolated A;Reference number: S30348; MUID:93238739; PMID:8097463 A;Accession: S30348
                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Reference number: PH0891; MUID:92078857; PMID:1836012
A;Accession: PH0924
A;Accession: PH0924
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PH0924
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A;Residues: 1-11 <GOL>
A;Cross-references: UNIPARC:UPI000017C9F3
A;Experimental source: concanavalin A-activated lymphoblast C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T-cell receptor beta chain V-D-J region (isolate 10) - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997 C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997 C;Accession: PH0924
C;Accession: PH0924
R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clotting protein - signal crayfish
C;Species: Pacifastacus leniusculus (signal crayfish)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P81070; UNIPROT:Q9UAR3; UNIPARC:UPI000017BDEA
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A;Molecule type: protein
A;Residues: 1-10 <KOP>
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Matches 3
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Pred. No. 2.8e+03;
1; Mismatches 0;
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potassium channel protein Slo G3 - fruit fly (Drosophila melanogaster) (fragment)
C;Species: Drosophila melanogaster
C;Species: Drosophila melanogaster
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Feb-1997
C;Accession: PS0443
R,Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta, A.; Bond Neuron 9, 209-216, 1992
A,Fitle: Calcium-activated potassium channels expressed from cloned complementary DNAs.
A;Reference number: JH0697; MUID:92360298; PMID:1497890
A;Accession: PS0443
A;Status: nucleic acid sequence not shown
A;Molecule type: mENA
A;Residues: 1-13 cADE>
A;Cross-references: UNIPARC:UPI000017BEBA
C;Comment: This potassium channel is activated by calcium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biochim. Biophys. Acta 1158, 227-236, 1993
A;Title: Characterization of two P-450 isozymes placed in the rat CYP2D subfamily. A;Reference number: S39761; MUID:94072607; PMID:8251521
A;Accession: S39762
A;Molecule type: protein
A;Residues: 1-12 <OHI>A;Cross-references: UNIPARC:UPI000017C8F7
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A;Cross-references: FlyBase:FBgn0003429
C;Keywords: alternative splicing; ion channel; potassium channel; transmembrane protein
Search completed: April
Job time : 20 secs
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Matches 3; Conservative
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streptomyce clivia mini rhizobium l

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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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              2006, 16:51:26; Search time 116.333 Seconds (without alignments) 54.582 Million cell updates
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(c) 1993 - 2006 Biocceleration
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Q6LCH4 RAT
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   RESULT 2
Q6LCH4 RAT PRELIMINARY;
ID Q6LCH4;
AC Q6LCH4;
AC Q6LCH4;
DT 05-JUL-2004 (TrEMBLrel. 27, Cr
DT 05-JUL-2004 (TrEMBLrel. 27, La
DT 05-JUL-2004 (TrEMBLrel. 27, La
DT 05-JUL-2004 (TrEMBLrel. 27, La
DE Glucose-6-phosphatase (Fragmen
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata;
OC Mammalia; Eutheria; Euarchontc
OC Muridae; Murinae; Rattus.
OX MCBI_TaxID=10116;
RN [1]
RN [1]
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SEQUENCE.
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QôLAP7 HUMAN
QôLAP7 HUMAN
PRELIMINARY;
AC
QôLAP7;
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Qò-JUL-2004 (TrEMBLrel. 27, La
DT
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RAMMBAIla; EMCHORICE
RY
MCLEOTIDE SEQUENCE
RY
PONGPHATABE Gene: transcription
RY
Glucocorticoids in H411E hepat
RY
PONGPHATABE GENERCE
RA
Allan Brian B;
RH
SUDMMITER
SQ
SEQUENCE
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AN; 1379 MW; AN
SEQUENCE
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AN; 1379 MW; AN
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Matches 4
                                            Glucose-6-phosphatase (Fragment).
Rattus norvegicus (Rat);
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognal
Muridae; Murinae; Rattus.
NCBI_TaxID=10116;
[1]
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STRAIN=Sprague-Dawley; TISSUE=Liver;
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Submitted (MAR-1996) to the EMBL/GenBank/DDBJ
EMBL; X96937; CAA65638.1; -; Genomic_DNA.
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SEQUENCE 12 AA; 1379 MW; A6B78EEFDE32C446
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MEDLINE=96.184388; PubMed=8612793; DOI=10.1016/0014-5793(96)00224-4; Schmoll D., Allan B.B., Burchell A.; Schmoll D., Allan B.B., Burchell A.; "Cloning and sequencing of the 5 region of the human glucose-6-phosphatase gene: transcriptional regulation by cAMP, insulin and glucocorticoids in H411E hepatoma cells."; FEBS Lett. 383:63-66(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Glucose-6-phosphatase (Fragment).
Homo sapiens (Human).
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Q9UC60_HUMAN
Q7M3S5_9TRYP
Q9UWM5_SULSO
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Q7TE19 9CLOS
Q7TE20 9CLOS
Q7TE23 9CLOS
Q7TE26 9CLOS
Q7TE26 9CLOS
Q7TE30 9CLOS
Q7TE30 9CLOS
Q957T7 9PERC
SODN STRGR
Q8HR43 CLIMI
Q52840 RHILO
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Last annotation update)
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Score

Match

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Minimum Maximum

DB DB

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Searched:

Scoring table:

Title: Perfect score:

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April

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RESULT 3
O95179 HUMAN
O95179;
ID O95179;
AC O95179;
AC O95179;
DT O1-MAY-1999 (TrEMBLrel. 10,
DT O1-MAY-1999 (TrEMBLrel. 11,
DT O1-DEC-2001 (TrEMBLrel. 19,
DT O1-DEC-2001 (TrEMBLrel. 19,
DT O1-DEC-2001 (TrEMBLrel. 19,
DE Glucose-6-phosphatase hydro
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordat
OC Mammalia; Eutheria; Euarcho
OC Homo.
OX NCBI_TaxID=9606;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RX MEDLINE=99156747; PubMed=10
RA Schmoll D., Wasner C., Hind
RA Burchell A.;
RT "Identification of a cAMP r
Thosphatase hydrolytic subu
RT transcriptional regulation
RT hepatoma cells.";
RL Biochem. J. 338:457-463 (199
DR EMBL; APO5135; AAD11621.1;
FT NON TER

14
SQ SEQUENCE 14 AA; 1666 FM;
       RESULT 4
Q86715 HUMAN
ID Q86715 H
AC Q86716;
DT 01-JUN-2
DT 01-JUN-2
DT 01-JUN-2
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DE FLJ20097
OS Homo sap
OC Eukaryot
OC Homo.
OX NCBI_Tax
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               flanking sequence.";
Diabetes 45:1563-1571 (1996).
EMBL; U57552; AAB19044.1; -;
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Glucose-6-phosphatase hydrolytic subunit (Fragment).
Homo sapiens (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Identification of a cAMP response element within the glucose-
phosphatase hydrolytic subunit gene promoter which is involved
transcriptional regulation by cAMP and glucocorticoids in H4IIF
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MEDLINE=99156747; PubMed=10024523;
Schmoll D., Wasner C., Hinds C.J.,
TISSUE=Peripheral Nervous System;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
MEDLINE=Peripheral Nervous System;
                                                                                                                                                                                     01-JUN-2003 (TrEMBLrel. 24,
01-JUN-2003 (TrEMBLrel. 24,
01-JUN-2003 (TrEMBLrel. 24,
FLJ20097 protein.
                                                                                                                                    Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                               Q86TJ6_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcriptional regulation by hepatoma cells.";
Biochem. J. 338:457-463(1999).
                                                                                             NCBI_TaxID=9606
                                                                 NUCLEOTIDE SEQUENCE.
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100.0%; F1
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Pred. No. 3.5e+03;
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Allan B.B., Walther R.,
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RESULT 5
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ID Q9TW74;
AC Q9TW74;
DT 01-MAY-2
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Q9TWT4;
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Q9TWT4;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-MAY-2000 (TrEMBLrel. 14, Last annotation update)
Q1-JUN-2000 (TrEMBLrel. 14, Last annotation update)
Hemoglobin (Fragment).
Lumbricus terrestris (Common earthworm).
Lumbricus terrestris (Common earthworm).
Lumbricus terrestris (Lumbricus).
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Submitted (MAR-2003) to the
EMBL; BC047757; AAH47757.1;
SEQUENCE 15 AA; 1843 MW;
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MEDLINE=93293879; PubMed=8514787;

MEDLINE=93293879; PubMed=8514787;

Ownby D.W., Zhu H., Schneider K., Beavis R.C., C
Ownby D.W., Zhu H., Schneider K., Beavis R.C., C
The extracellular hemoglobin of the earthworm,
                                                                                                                                                                                                                                                                                                                                              Determination of subunit stoichiometry.";
J. Biol. Chem. 268:13539-13547(1993).
SEQUENCE 15 AA; 1665 MW; F9B83FE0050D
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EMBL; U07130; AAA74412.1; -;
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Q66874;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE. STRAIN=255;
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SEQUENCE 11 AA; 1330 MW
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-I- MISCELLANEOUS: On the 2D-gel the determined pI of the protein is: 5.5, its MW is: 48 kDa.
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TISSUE=Fibroblast;
MEDLINE=95009907; PubMed=7523108;
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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Pred. No.
1; Mismatc
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Pred. No.
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RESULT 9
Q7M063 MOUSE
ID Q7M063;
AC Q7M063;
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Calnexin (Fragment).
OS Mus musculus (Mouse).
OC ENKARYOCE; Metazoa; Chordat;
OC Muridae; Murinae; Mus.
OX NCBI_TaxID=1090;
RN [1]
RN MEDLINE=94216347; PubMed=81
RX MEDLINE=94416347; PubMed=81
RX MEDLINE=94416347; PubMed=81
RX MEDLI
      RESULT 10
FIBB CERSI
ID FIBB C
AC P14537
DT 01-JAN
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Best Local S
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   FIBB_CERSI S'
P14537;
01-JAN-1990 (Rel
01-JAN-1990 (Rel
01-FEB-2005 (Rel
Fibrinogen beta
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01-NOV-1996 (TrEMBLr
01-NOV-1996 (TrEMBLr
01-NOV-1998 (TrEMBLr
Polymerase (Fragment
Feline calicivirus
Fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIINE=94216347; PubMed=8163531;
Lenter M., Vestweber D.;
Lenter M., Vestweber D.;
"The integrin chains beta-1 and alpha-6 associate Calmexin prior to integrin assembly.";
J. Biol. Chem. 269:12263-12268(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Glires; Rodentia; Sciurogna
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON TER
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      (Rel. 13, (Rel. 46, beta chain
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                                                                                                   (Rel.
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Last sequence update)
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[Contains: Fibrinopeptide
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Pred. No. 4.2e+03;
1; Mismatches 2
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Matches 3
SEQUENCE
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01-NOV-1996 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                         Q47600 ECOLI PRELIMINARY;
Q47600;
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PROSITE; PS00514; FIBRIN AG C DOMAIN; PARTIAL.
Blood coagulation; Coiled coil; Direct protein
1 10 Fibrinopeptide B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROTEIN SEQUENCE
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                             "A family of regulatory genes associated with modification systems.";
J. Bacteriol. 173:1367-1375(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                    Enterobacteriaceae;
NCBI_TaxID=562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NON TER
                                                                                                 NUCLEOTIDE SEQUENCE.
MEDLINE=91139577; PubMed=1995588;
                                                                                                                                                                Bacteria; Proteobacteria;
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                                                                                                                                                                                                      REase protein (Fragment)
                                                                                     Tao T.
                                                                                                                                                                                                                                                                                   ECOLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aggregation.
SUBUNIT: Heterohexamer; disulfide linked. Contains 2 sets of 3
SUBUNIT: Heterohexamer; disulfide linked. Contains 2 sets of 3
nonidentical chains (alpha, beta and gamma). The 2 heterotrimers
are in head to head conformation with the N-termini in a small
central domain (By similarity).

COMAIN: A long coiled coil structure formed by 3 polypeptide
chains connects the central nodule to the C-terminal domains
(distal nodules). The long C-terminal ends of the alpha chains
fold back, contributing a fourth strand to the coiled coil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: Fibrinogen has a polymerize into fibrin and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 structure.
                     M63619; AAA24556.1;
ER 11 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : Conversion of fibrinogen to fibrin is triggered by thrombin, ch cleaves fibrinopeptides A and B from alpha and beta chains, thus exposes the N-terminal polymerization sites responsible the formation of the soft clot.
                                                                                                                                                                                                                                                                                                                                                                                       ω,
                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                     Bourne J.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          1097 MW;
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                                                                                     , Blumenthal R.M.;
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100.0%;
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19,
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          63175479572AB5A4 CRC64;
                                   Genomic_DNA
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RESULT 13

QPR5D6_CHRVI
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DT 01-MAY-2
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QFR5D6_CHRVI PRELIMINARY;
QFR5D6;
01-MAY-2000 (TrEMBLrel. 13, 0
01-MAY-2000 (TrEMBLrel. 13, I
01-JUN-2000 (TrEMBLrel. 14, I
POLY(3-HYDROXYBUTYRIC acid) s
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Biochem.
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STRVN
XYLA_S
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Streptomyces violaceoruber. Bacteria; Actinobacteria; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JAN-1990
01-JAN-1990
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                                                                                                                                                                                                                                               InterPro; IPR001998; XyLose isom.
InterPro; IPR001998; XyLose isom.
PROSITE; PS00172; XYLOSE ISOMERASE 1; PA:
Carbohydrate metabolism; Direct potein
Magnesium; Metal-binding; Pentose shunt;
Magnesium; Metal-binding; Pentose shunt;
                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse as long as its content is in no way modified and this s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=90104230;
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                                                                                                                                                                                                                                                                                                                                                                                                      SUBUNIT: Homotetramer.
SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the xylose isomerase family.
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LYTIC ACTIVITY: D-xylose = D-xylulose
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13, Last sequence update)
47, Last annotation update)
(EC 5.3.1.5) (Fragment).
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                                                                                                                                                                                                        1376 MW;
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granule-associated 41 kDa
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Pred. No. 6.
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D-xylose isomerase from Stro
chemical derivatization and
                                                                                                                                                                                                                                                             shunt;
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7.2e+03;
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RESULT 14
Q778X5 PLAYO PRELIMINARY;
AC Q778X5;
AC Q778X6;
AC Q778X12 Alveolata;
Apicon
AN Ame=PY07095;
Alsemodium yoelii yoelii.
OC Eukaryota; Alveolata; Apicon
OX NCBL_TaxID=73239;
AN QCLEOTIDE SEQUENCE.
AN AMEDLINE=22255706; PubMed=122
AN AMEDLINE=2255706; PubMed=122
AN AMEDLINE=22255706; PubMed=122
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RESULT 15
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Matches 4
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MEDLINE=22255706; pubMed=12368865; DOI=10.1038/nature01099;
Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M.,
Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L.,
Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L.,
Shallom S.J., van Aken S.E., Riedmiller S.B., Felbblyum T.V.,
Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,
Florens L., Yates J.R. III, Raine J.D., Sinden R.E., Harris M.A.,
Cunningham D.A., Preiser P.R., Bergman L.W., Vaidya A.B.,
van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R.,
Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Genome sequence and comparative analysis parasite Plasmodium yoelii yoelii."; Nature 419:512-519(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEMS Microbiol. Lett. 78:227-232(1992). SEQUENCE 15 AA; 1874 MW; 165FA9A16E
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MEDLINE-93146381; PubMed=1490603; DOI=10.1016/0378-1097(92)90031-I;

Liebergesell M., Schmidt B., Steinbuchel A.;

"Isolation and identification of granule-associated proteins relevant for poly(3-hydroxyalkanoic acid) biosynthesis in Chromatium vinosum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacteria; Proteobacteria; Gammaproteobacteria; Chromatiales; Chromatiaceae; Allochromatium.
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EMBL; J01670; AAA24475.1; -; Genomic_DNA.
NON TER 9
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Escherichia coli Kl2.
Bacteria; Proteobacteria;
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                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
MEDLINE=83195078; PubMed=6302686;
Medura T.A., Roof W.D., Foltermann K.F.,
                                                                                                                                                                                                                                                                                          pyrB-pyrI operon."
Proc. Natl. Acad. '
                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
MEDLINE=82275057; PubMed=7051000;
Pauza C.D., Karels M.J., Navre M., Schachman H.K.;
"Genes encoding Escherichia coli aspartate transcarbamoylase:
                                                                                                                                                                                                                                                                                                                                                                        Enterobacteriaceae; Escherichia.
NCBI_TaxID=83333;
                                                                                                                                                                                                        catalytic
                                                                                                                                                                                                      "Nucleotide sequence of the structural catalytic polypeptide of aspartate tran
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completed: April
ne : 117.333 secs
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              2006, 17:02:58
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                                                                                                                                              99EFD723344AA1F1 CRC64;
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                                                                                                         Score 19;
Pred. No.
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nscarbamoylase of
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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  seq length: 0
seq length: 15
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1: /cgm2 6/ptodata/1,
2: /cgm2 6/ptodata/1,
3: /cgm2 6/ptodata/1,
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Match Length
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/cgm2_6/ptcdata/1/iaa/H_COMB.pep:*
/cgm2_6/ptcdata/1/iaa/ETUS_COMB.pep:*
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US-09-489-760-14
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US-09-489-760-14
US-09-269-250E-26
US-09-489-760-4
US-09-489-760-4
US-09-489-760-3
US-09-489-760-3
US-09-803-126-5
US-09-803-126-5
US-09-803-916-5
US-09-803-917-5
US-08-724-648-21
US-08-724-548-21
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US-09-269-250E-20
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GENERAL INFORMATION:
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ALIGNMENTS

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FRUCIL NO. 100.03.

FREEZEL NO. 1 INFORMATION:

APPLICANT: Rijksuniversiteit Te Leidin

APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Hunt, Donald F

APPLICANT: Hard, Victor H

ITITLE OF INVENTION: The HA-1 Antigen

FILE REFERENCE: 2183-4285US

CURRENT FILING DATE: 2000-01-21

EARLIER APPLICATION NUMBER: PCT/NL98/00424

EARLIER APPLICATION NUMBER: PCT/NL98/00424

EARLIER FILING DATE: 1998-07-23

NUMBER OF SEQ ID NOS: 16

SOFTWARE: PATENTIN Ver. 2.1

SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 2
US-09-489-760-2
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CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
LENGTH: 9
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Best Local Similarity
Matches 9; Conserv
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Histocompatibility antigen
                        TYPE:
                                           ENGTH: 9
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Pred. No. 4.6e+05;
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US-09-269-250E-28
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                                                                                                                          US-09-489-760-16
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APPLICANT: Hunt, Denald F
APPLICANT: Hunt, Denald F
APPLICANT: Hunt, Denald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: GOLINY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 28
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Best Local S
Matches 9
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SEQ ID NO 16
LENGTH: 13
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GENERAL INFORMATION:
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Best Local
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ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: PCR Product
-09-269-250E-28
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GENERAL INFORMATION:
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OTHER INFORMATION: Description
                                                                                                                                                                     ORGANISM: Unknown Organism
                                                                                                                                                                                     TYPE: PRT
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VLHDDLLEA 11
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                                                         Score 45; DB 2;
Pred. No. 0.034;
Mismatches 0
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Pred. No. 0.034;
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US-09-489-760-5
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US-09-269-250E-18
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US-09-269-250E-26
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CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 5
Sequence 26, Application US/09269250E
PACENT NO. 6830883
GENERAL INFORMATION:
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
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APPLICANT: GOULTMY, RISHA
APPLICANT: GOULTMY, RISHA
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT FILLYG DATE: 1999-05-21
CURRENT FILLYG DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
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Patent No. 6830883
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Best Local Similarity
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ORGANISM:
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OTHER INFORMATION: Descriptio
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TYPE: PRT
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Pred. No. 4.6e
0; Mismatches
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Pred. No. 4.6e+05;
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GENERAL INFORMATION:

APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: DET/NL98/00424
EARLIER APPLICATION WUMBER: PCT/NL98/00424
EARLIER APPLICATION WIMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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US-09-269-250E-29
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Best Local Similarity
Thes 8; Conserve
US-09-269-250E-29
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US-09-489-760-14
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                                                                                                                                                  Sequence 29, Application US/09269250E
Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
         LENGTH: 9
TYPE: PRT
ORGANISM: Human
FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a
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SEQ ID NO 14
LENGTH: 13
                                                                                                                                     SEQ ID NO 29
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Unknown Organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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3 VLRDDLLEA 11
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Pred. No. 1.1;
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Sequence 4, Application US/09489760
Patent No. 6878375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
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                                                                                                                                   SEQ ID NO 4
LENGTH: 9
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Best Local Similarity
Matches 8; Conserv
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Patent No. 6878375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                       FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: The HA-1 Antigen FILE REFERENCE: 2183-4285US CURRENT APPLICATION NUMBER: US/09/489,760 CURRENT FILING DATE: 2000-01-21 CURRENT FILING DATE: 2000-01-21 EARLIER FILING DATE: 1998-07-23 EARLIER FILING DATE: 1998-07-23 NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
ORGANISM: Histocompatibility antigen
FEATURE:
FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (2)..(7)
OTHER INFORMATION: AMINO ACIDS X REPRESENT LEUCINE OR ISOLEUCINE
OTHER INFORMATION: RESIDUES
                                                                                                                    TYPE: PRT
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NAME/KEY: PEPTIDE
LOCATION: (3)
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TYPE: PRT
ORGANISM: Histocompatibility antigen
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Pred. No. 4.6e+05;
0; Mismatches 1
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Pred. No. 4.6e+05;
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US-09-269-250E-38
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US-09-025-819-5
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Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: GOULMY, E18a
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT PILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
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Best Local Similarity 80...
Thes 8; Conservative
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Patent No. 6225097
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: UP 251675
PILING DATE: 17-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
REFERENCE/DOCKET NUMBER: 10235/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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STATE: USA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Obata, Shusei
APPLICANT: Nishino, Tokuzo
APPLICANT: Koyama, Tanetoshi
APPLICANT: Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 8
                                                                                                                                                                                                                 ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0:
FILING DATE: 19-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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1500 K Street, N.W.
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Pred. No. 4.6e+05;
0; Mismatches 3
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       10235/2
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Pred. No. 4.6e+05;
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                                                       ; TOPOLOGY: linear; WOLECULE TYPE: peptide; SEQUENCE DESCRIPTION: SEQ ID NO: 5: US-09-808-126-5
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US-09-808-126-5
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                                                                                                                                           COUNTRITUDE.

COUNTRITUDE.

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Vers

APPLICATION NUMBER: US/09/808,126

FILING DATE: 08-May-2001

CLASSIFICATION : <Unknown>
PRIOR APPLICATION NUMBER: 09/025,819

FILING DATE: <Inknown>
PRIOR APPLICATION NUMBER: 39,546

REFERENCE/DOCKET NUMBER: 10235/2

TELEFONE: 20-220-4200

TELEFONE: 20-220-4201

INFORMATION FOR SEQ ID NO: 5:

SEQUENCE CHARACTERISTICS:

LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/09808126
Patent No. 6410280
GENERAL INFORMATION:
APPLICANT: Obata, Shusei
Query Match
Best Local Similarity
Matches 4; Conserv
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
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Local Similarity 80.0%;
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Koyama, Tanetoshi
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Sato, Yoshihiro
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Score 25; DB 2; ~~
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RESULT 15

US-09-801-51-5

Jacquence 5. Application US/09803951

Patent No. 6413761

Patent No. 6413761

Jacquence 5. Application US/09803951

Jacquence 5. Application US/09803951

Jacquence 5. Application Jacquence 1

Jacquence 7. Jacquence 1

Jacquence 7
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Maximum Match 100%
Listing first 45 summaries
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1: /cgm2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgm2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgm2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgm2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

5: /cgm2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*

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US-10-81-335-1
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US-10-623-176-45
US-10-623-176-45
US-10-623-176-10
US-10-623-176-10
US-10-623-176-10
US-10-623-176-10
US-10-623-176-47
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US-10-791-217-1
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US-10-791-217-1
US-10-791-217-1
US-10-791-217-1
US-11-007-740-26
US-10-623-176-40
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US-10-623-176-6-60
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Sequence 2, Appli
Sequence 1, Appli
Sequence 20, Appli
Sequence 43, Appl
Sequence 16, Appli
Sequence 16, Appli
Sequence 16, Appli
Sequence 41, Appli
Sequence 45, Appli
Sequence 10, Appli
Sequence 5, Appli
Sequence 14, Appli
Sequence 16, Appli
Sequence 20, Appli
Sequence 21, Appli
Sequence 22, Appli
Sequence 24, Appli
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RESULT 1 US-10-623-176-2 ; Sequence 2, A ; Sequence 2, A ; Publication N GENERAL INFOR APPLICANT: G APPLICANT: ; APPLICANT: ; TITLE OF INFOR FILE REFEREN CURRENT APPL		45	44	43	42	41	40	39	38	37	36	35	34	ယ္	32	31	30	29	28	
SULT 1 -10-623-176-2 -10-623-176-2 Sequence 2, Applicat Publication No. US20 GENERAL INFORMATION: APPLICANT: GUNT, APPLICANT: Hunt, C APPLICANT: Engelha TITLE OF INVENTION: FILE REFERENCE :18 FILE REFERENCE :18 CURRENT APPLICATION DATE PRIOR APPLICATION A		27	27	27	28	28	28	28	28	28	28.5	28.5	29	30	30	30	32	.32	33	
SULT 1 -10-623-176-2 -10-623-176-2 Sequence 2, Application US/10623176 Sequence 2, Application US/10623176 Sequence 2, Application US/200446A1 GENERAL INFORMATION: APPLICANT: HUNT, Donald F. APPLICANT: Engelhard, Victor H. TITLE OF INVENTION: HA-1 epitopes an FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/62 CURRENT FILING DATE: 2003-07-18 PRIOR APPLICATION NUMBER: 09/489,760			60.0	•	•	•	•	•	•	62.2	•	u	64.4	66.7	66.7	•	71.1	71.1		
ttion US/1 004009244 : Els A.J. Bonald r. ard, Vict : HA-1 ep 83-6047US 81-2003- E: 2003-		10	10	9	13	10	9	9	9	9	œ	80	9	9	9	ø	14	9	9	
M. Alloca		4	4.	S	4	4	4	4	4	4.	თ	4	4	σ	4	4,	4	4	4.	
ULT 1 10-623-176-2 equence 2, Application US/10623176 ublication No. US20040092446A1 ublication No. US20040092446A1 ENERAL INFORMATION: APPLICANT: Goulmy, Els A.J.M. APPLICANT: Hunt, Donald F. APPLICANT: Engelhard, Victor H. TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/623,176 CURRENT FILING DATE: 2003-07-18 PRIOR APPLICATION NUMBER: 09/489,760	ALIGNMENTS	US-10-702-400-90	-10-623-176-8	-10-705-459-2	US-10-623-176-64	-10-623	-10-837-217-9	-10-777-	-10-623-176-3	US-10-623-176-23	-11-007-740-3	US-10-623-176-15	US-10-623-176-48	US-11-007-740-40	US-10-791-217-4	US-10-623-176-14	-10-623-	US-10-623-176-4	US-10-623-176-46	
		90	8		64	7,	95	Sequence 95,						40	4,	Sequence 14,	50	Sequence 4,	46	
		App1	Appli	280, App	Appl	Appli	App1	App1	Appl	Appl	Appl	Appl	Appl	, Appl	•		, Appl	Appli	App1	

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SEC 12.

LENGTH: 9

ITYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE: PRT
OTHER INFORMATION: Description
FEATURE:
NAME/KEY: SITE
LOCATION: (1)..(9)
US-10-623-176-2

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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR PILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTLIN VEY: 2.1
SOFTWARE: PATENTLIN VEY: 2.1
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Best Local S
Matches 9
Sequence 2, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
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Similarity 100.0%;
9; Conservative 0
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                                                                                                                                                                                                         RESULT 4
US-11-007-740-20
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                    Sequence 20, Application US/11007740

Publication No. US20050233350A1

GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
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SEQ ID NO 2
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial sequence
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Publication No. US20050031612A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/861,335
CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: PCT/NIO2/00791
PRIOR FILING DATE: 2002-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Goulmy, Elsa A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot TITLE OF INVENTION: of tumors
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CURRENT FILING DATE: 2004-03-02
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PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
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PRIOR FILING DATE: 1999-05-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: HA-1 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 45; DB 5;
100.0%; Pred. No. 1.7e+06
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          FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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US-10-623-176-76
Sequence 76, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Bigelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-11-007-740-20
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PRIOR FILING DATE: 2000-01-21
PRIOR PPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: 07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                  RESULT 6
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Publication No. US20040092446A1
GENERAL INFORMATION:
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
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ORGANISM: Artificial Sequence
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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TYPE: PRT
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100.0%; Pred. No. 0.17;
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Pred. No. 1.7e+06;
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PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEO ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 16
ENOTH: 13
RESULT 8
US-11-007-740-28
; Sequence 28, Application US/11007740
; Publication No. US20050233350A1
; GENERAL INFORMATION:
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Best Local Similarity
Thehes 9; Conserv:
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; LOCATION: (1)..(13)
US-10-623-176-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16, Application US/10791217 Publication No. US20040191268A1 GENERAL INFORMATION:
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Best Local (
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 76
                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Goulny, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 2183-4285US
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PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
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                                                                                                                                                                                                                                                                                      OTHER INFORMATION: obtained from HA-1+/+ phenotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: KIAA0223
OTHER INFORMATION: sequence derived from a presumed HA-1 homozygous
OTHER INFORMATION: positive individual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
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Pred. No. 0.22;
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RESULT 9

US-10-623-176-41
; Sequence 41, Application US/10623176
; Sublication No. US20040092446A1
; GENERAL INFORMATION:
; APPLICANT: Goulny, Els A.J.M.
; APPLICANT: Hunt, Donald F.
; APPLICANT: Engelhard, Victor H.
; TITLE OF INVENTION: HA-1 epitopes and uses thereof
; TITLE OF EFFRENCE: 2183-6047US
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                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR PILING DATE: 2000-01-21

PRIOR PILING DATE: 2000-01-21

PRIOR FILING DATE: 1997-07-23

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR FILING DATE: 1998-07-23

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR FILING DATE: 2000-01-24
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SEQ ID NO 28
LENGTH: 13
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                                                                                                    Query Match
Best Local
                                                                                   Matches
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Best Local (
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                                                                                                                                                                           NAME/KEY: SITE LOCATION: (1)..(9)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                       FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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                         1 VLHDDLLE 8
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                                                                               Similarity
8; Conserv
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VLHDDLLE
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                                                                               91.1%; Score 41; DB
100.0%; Pred. No. 1.7
tive 0; Mismatches
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100.0%; Pred. No. 0.22;
tive 0; Mismatches
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                                                                                                                     Length 9;
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RESULT 10
US-10-623-176-45
; Sequence 45, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:

Hunt, Donald F.

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GENERAL INFORMATION:

APPLICANT: Goulmy, Els A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

TITLE OF INVENTION: HA-1 epitopes and uses thereof

FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR FILING DATE: 2000-01-21

PRIOR FILING DATE: 1997-07-23

PRIOR FILING DATE: 1997-07-23

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR FILING DATE: 1998-07-23

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR FILING DATE: 2000-01-24

NUMBER OF SEQ ID NOS: 77

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 10

LENGTH: 9
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PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 2000-01-21
PRIOR PILING DATE: 1997-07-23
PRIOR PELICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR PILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR PILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN VOI. 2.1
SEQ ID NO 45
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Query Match
Best Local Similarity
"atches 8; Conserva
                                                                                                                US-10-623-176-10
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CURRENT FILING DATE: 2003-07-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: SITE
LOCATION: (1)..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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                                                                                                                               NAME/KEY: SITE
LOCATION: (1)..(9)
                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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                                                                                                                                                                                FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
nes 8; Conserva
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                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91.1%; Score 41; DB 4; 100.0%; Pred. No. 1.7e+(
                                          82.2%;
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    Mismatches

                                          Score 37; DB 4; Length 9; Pred. No. 1.7e+06;
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APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 1900-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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US-10-623-176-47
; Sequence 47, Application US/10623176
; Publication No. US20040092446A1
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LOCATION: (1)..(9)
US-10-623-176-47
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US-10-791-217-5
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Best Local Similarity luv.
Thes 7; Conservative
                                                                                                                                                                                                                                       PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Goulimy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
APPLICANT: Enge
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SEQ ID NO 47
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                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
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TYPE: PRT
ORGANISM: Artificial Sequence
OTHER INFORMATION: obtained from KIAA0223 partial complementary DNA
                                                                                                  TYPE: PRT ORGANISM: Artificial sequence
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OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                           FEATURE:
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100.0%; Pred. No. 1.7e+06;
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RESULT 14

US-11-007-740-18

Sequence 18, Application US/11007740

Publication No. US20050233350A1

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa

ITITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1

FILE REFERENCE: 2799/58994-A

CURRENT APPLICATION NUMBER: US/11/007,740

CURRENT FILING DATE: 2004-12-08

PRIOR APPLICATION NUMBER: 09/269,250

PRIOR APPLICATION NUMBER: 09/269,250

PRIOR FILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 40

SOPTWARE: Patentin version 3.1

SEQ ID NO 18

LENGTH: 9

TYPE: PRT

ORGANISM: Artificial Sequence
; OTHER INFORMATION: D. FEATURE: ; PEATURE: ; NAME/KEY: SITE ; LOCATION: (1)..(10) US-10-623-176-44
                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1997-07-23
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: D2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOPTWARE: PATENTIN VET: 2.1
SEQ ID NO 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 15
US-10-623-176-44
US-10-623-176-44
; Sequence 44, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
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Best Local Similarity
Watches 8; Conserve
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                  LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                FEATURE:
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                                                                        Description of Artificial Sequence: HA-1 peptide
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Pred. No. 1.7e+06;
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Query Match

Best Local Similarity

Best Local No. 4.4;

Matches 1; Indels 0; Gaps

Best Local No. 4.4;

Best Local No
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Result
No.
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Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Database
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Maximum Match 10
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     seq length: 0 seq length: 15
        Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/US06_NI

2: /SIDSS/ptodata/2/pubpaa/US06_NI

3: /SIDSS/ptodata/2/pubpaa/US07_NI

4: /SIDSS/ptodata/2/pubpaa/US09_NI

6: /SIDSS/ptodata/2/pubpaa/US10_NI

7: /SIDSS/ptodata/2/pubpaa/US11_NI

8: /SIDSS/ptodata/2/pubpaa/US06_NI
      US-10-791-217A-2
45
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/SIDS5/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US17_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US17_NEW_PUB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                               GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100%
                      US-11-010-748A-11
US-11-010-748A-12
US-11-129-143-163
US-11-129-143-165
US-11-129-143-166
US-11-129-143-166
US-11-129-143-166
US-11-129-143-166
US-11-129-143-162
US-11-004-399-1155
US-10-485-788A-353
US-10-485-788A-353
US-10-485-788A-355
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US-11-033-039-809
US-11-033-039-809
US-11-033-039-809
US-11-004-399-3188
US-11-004-399-3188
US-11-004-399-3210
US-11-166-288-11
US-11-166-288-11
US-11-166-288-11
US-11-168-7435A-443
US-10-885-7435A-443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUMMARIES
Sequence 11, Appl Sequence 163, App Sequence 164, App Sequence 165, App Sequence 161, App Sequence 161, App Sequence 115, App Sequence 115, App Sequence 115, App Sequence 354, App Sequence 354, App Sequence 354, App Sequence 354, App Sequence 357, App Sequence 358, App Sequence 809, App Sequence 809, App Sequence 801, App Sequence 821, App Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 413, App Sequence 413, App Sequence 413, App Sequence 413, App Sequence 414, App Sequence 4144, App Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Description
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ALIGNMENTS

US-11-010-748A-11

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Sequence 11, Application US/11010748A

Publication No. US20050244421A1

GENERAL INFORMATION:
APPLICANT: MATCH MACKE PALENT GMDH
APPLICANT: STRITTMATTER, Wolfgang
APPLICANT: SCHARM, Burkhard
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE
FILE REFERENCE: MER-136
CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
PRIOR APPLICATION NUMBER: EPTS/EP03/06251
PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EPTS/EP03/06251
PRIOR PILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
SOPTMARE: PatentIn version 3.1
SEQ ID NO 11
LENGTH: 9
                                                                                                                                                   RESULT 2
US-11-010-748A-12
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Sequence 12, Application US/11010748A
Publication WO. US20050244421A1
GENERAL INFORMATION:
APPLICANT: Merck Patent GmbH
APPLICANT: STRITTMATTER, Wolfgang
APPLICANT: WOLL, Heidrun
APPLICANT: SCHARM, Burkhard
                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (1)...(9)
OTHER INFORMATION: ell epitopes
                                                                                                                                                                                                                                                                                                                     y Match 100.0%; Score 45; DB 7;
Local Similarity 100.0%; Pred. No. 1.4e+05;
hes 9; Conservative 0; Mismatches 0;
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, NAME/KEY: MISC FEATURE

, LOCATION: (1)...(9)

, OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T

, OTHER INFORMATION: ell epitopes

US-11-010-748A-12
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US-11-129-143-163
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                                                                                                   Sequence 164, Application US/11129143 Publication No. US20050266518A1 GENERAL INFORMATION:
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SEQ ID NO 163
LENGTH: 6
TYPE: PRT
ORGANISM: Bacillus stearothermophilus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 163, Application US/11129143 Publication No. US20050266518A1
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Best Local Similarity
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PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                         -11-129-143-163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
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CURRENT FILING DATE: 2004-12,13
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                                                  APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 197
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TYPE: PRT
ORGANISM: Homo sapiens
                                  APPLICANT:
                                                                                   APPLICANT: BERRY, Alan
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APPLICANT:
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8; Conserv
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BRETZEL, Werner
HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
MAYER, Anne F.
YELISEEV, Alexei A.
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HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
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Pred. No.
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Pred. No. 1.4e+05;
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CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: PatentIn version 3.1
SEQ ID NO 166
LENGTH: 6

APPLICANT:
APPLICANT:
APPLICANT:

LOPEZ-ULIBARRI, Rual MAYER, Anne F.

APPLICANT: BERRY, Alan APPLICANT: BRETZEL, W

BRETZEL, Werner HUMBELIN, Markus

APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143

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                                                RESULT 6
US-11-129-143-166
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 164
Sequence 166, Application US/11129143
Publication No. US20050266518A1
GENERAL INFORMATION:
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Best Local :
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SEQ ID NO 165
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APPLICANT:
APPLICANT:
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APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, W
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                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Publication No.
                                                                                                                                                                                                                                                                                                                                       APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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TYPE: PRT
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Local Similarity 80.0%;
hes 4; Conservative
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                                                                                                                                              2 LHDDL 6
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HUMBELIN, Markus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/11129143
                                                                                                                                                                              Conservative
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                                                                                                                                                                                           57.8%;
80.0%;
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Pred. No. 1.4e+05;
1; Mismatches 0
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APPLICANT: BRETZEL, Werner

APPLICANT: HUMBELIN, Markus

APPLICANT: LOPEZ-ULIBARRI, Rual

APPLICANT: LOPEZ-ULIBARRI, Rual

APPLICANT: WAYER, Anne F.

APPLICANT: YELISEEV, Alexei A.

TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION

FILE REFERINCE: C38435/121966

CURRENT APPLICATION NUMBER: US/11/129,143

CURRENT APPLICATION NUMBER: US/11/129,143

CURRENT FILING DATE: 2005-05-13

NUMBER OF SEQ ID NOS: 197

SOFTWARE: Patentin version 3.1

SEQ ID NO 162

LENGTH: 6

TYPE: PRT
ORGANISM: Rhizobium sp. strain NGR234

US-11-129-143-162
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APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: WAYER, Anne F.
APPLICANT: YELLSEBV, Alexel A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION UNUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 161
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US-11-129-143-161
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US-11-129-143-161
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                                                                                              Matches
                                                                                                                     Query Match
Best Local 9
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APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: BERRY, Alan
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                                                                                                                     Similarity
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80.0%;
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1; Mismatches
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Pred. No.
                                                                                     Score 25; DB 7; Le
Pred. No. 1.4e+05;
1; Mismatches 0;
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RESULT 11
US-10-485-788A-353
; Sequence 353, Application US/10485788A
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                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide US-10-931-251A-13
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US-10-931-251A-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: PRT; ORGANISM: SARS-COV Virus US-11-004-399-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: KOLOKITHAS, ANGELO STEPHEN
APPLICANT: MONBLI, CRAIG ROBERT
TITLE OF INVENTION: PROTEIN KINASE AND PHOSPHATASE SUBSTRATES AND MULTIPLEX
TITLE OF INVENTION: ASSAYS FOR IDENTIFYING THEIR ACTIVITIES
FILE REFERENCE: STG-106
CURRENT APPLICATION NUMBER: US/10/931,251A
CURRENT FILING DATE: 2004-09-01
NUMBER OF SEQ ID NOS: 42
SOPTWARE: PATENTIN Ver. 3.2
SEQ ID NO 13
LENGTH: 15
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Best Local Similarity 100.
Conservative
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Publication No. US20066053516A1
GENERAL INFORMATION:
APPLICANT: Chye, Mee Lee
APPLICANT: Li, Hong Ye
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13, Application US/10931251A
Publication No. US20060046277A1
GENERAL INFORMATION:
APPLICANT: BELYAEV, ALEXANDER SERGEEVICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1 SEQ ID NO 1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ramalingam, Sathiskumar
APPLICANT: Poon, Leo Lit Man
APPLICANT: Peiris, Joseph Sriyal Malik
TITLE OF INVENTION: Genetically Modified Plants Comprising SARS-COV Viral Nucleotide
TITLE OF INVENTION: Sequences and Methods of Use Thereof For Immunization Against SA
FILE REFERENCE: 2587/73166/RDK
CURRENT APPLICATION NUMBER: US/11/004,399
CURRENT APPLICATION NUMBER: US 60/527,637
PRIOR APPLICATION NUMBER: US 60/527,637
PRIOR FILING DATE: 2003-12-03
NUMBER OF SEQ ID NOS: 4043
COPTEMBER OF SEQ ID NOS: 4043
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                          Score 24;
; Pred. No.
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TITLE OF INVENTION MOLECULAR INTERACTIONS in Cells
FILE REFERENCE: 20054-003320US
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR APPLICATION NUMBER: W0 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: W0 PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: Patentin version 3.1
SEQ ID NO 354
LENGTH: 7
TYPE: PRT
ORGANISM: Homo sapiens
RESULT 13
US-10-485-788A-355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
APPLICANT: Schweizer, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WG PCT/US02/24655
PRIOR APPLICATION NUMBER: WG PCT/US02/24655
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; Sequence 354, Application US/10485788A
; Publication No. US20050282743A1
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; ORGANISM: Homo sapiens
US-10-485-788A-353
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                                                                                                                                                Query Match
Best Local Similarity 50.0
""" Conservative
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Best Local Similarity 50.0%;
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz,
APPLICANT: Schweizer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Lu, Peter S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION:
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                                                                                             LVHDDV 7
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Arbor Vita Corporation
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Pred. No. 1.4e+05;
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Sequence 785, Application US/1103039

Publication No. US20060002947A1

GENERAL INFORMATION:
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN

TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01

CURRENT FILING DATE: 10/245,871
PRIOR APPLICATION NUMBER: US/11/033,039

CURRENT FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR APPLICATION NUMBER: 10/297,000
PRIOR FILING DATE: 2002-09-17
PRIOR FILING DATE: 2002-09-17
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/297,000
PRIOR FILING DATE: 1999-09-14
PRIOR APPLICATION NUMBER: 09/396,813
PRIOR RILING DATE: 1999-09-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR PILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR APPLICATION NUMBER: W0 PCT/US02/24655
PRIOR PILING DATE: 2002-02-25
PRIOR PILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
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US-11-033-039-785
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                                                                                                                                                                                                 US-11-033-039-785
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Best Local Similarity 50.0
Whiches 3; Conservative
                                                                                            Query Match 51.1%; Score 23; DB 7; Length 9; Best Local Similarity 44.4%; Pred. No. 1.4e+05; Matches 4; Conservative 3; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 355, Application US/10485788A Publication No. US20050282743A1
                                                                                                                                                                                                                                                                                                 SEQ ID NO 785
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APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 1452
                                                                                                                                                                                                                        ORGANISM: Zaire ebolavirus
                                                                                                                                                                                                                                                LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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Pred. No. 1.4e+05;
3; Mismatches 0;
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RESULT 15 US-11-033-039-809 ; Sequence 809, Application US/11033039 ; Publication No. US20060002947A1

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GENERAL INPORMATION:
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
TITLE OF INVESTION: L1-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT APPLICATION NUMBER: 10/245,871
PRIOR FILING DATE: 2002-09-71
PRIOR PILING DATE: 2002-09-71
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-09-71
PRIOR APPLICATION NUMBER: 109/396,813
PRIOR RILING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 1452
SOFTWARE: PATE ANTENING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 1452
SOFTWARE: PATE OF SEQ ID NOS: 1452
SOFTWARE: PATE OF SEQ ID NOS: 1452
SOFTWARE: DATE: 1999-09-14
DOBANISM: Zaire ebolavirus
US-11-033-039-809

QUESTY MATCH
Best Local Similarity 44.4%; Pred. No. 1.4e+05;
MATCHES 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;
MATCHES 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

Search completed: April 6, 2006, 17:29:30

Search completed: April 6, 2006, 17:29:30
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Maximum Match
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Maximum DB
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Perfect score:
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1: geneseqp1980
2: geneseqp2000
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my v odor 22/e2hw		B f	Human	Adt72766 Human RSV	Aau71428 Human MHC		Abp71119 E10 prote	Aag94234 Human com	Adt72767 Human RSV	7	-	Adt73023 Human RSV	T-cell	Aaw99195 Minor his	Aaw97415 KIAA0223	_	Adh40333 Human min	Aaw97375 HA-1 H-al	Aaw99196 Minor his	Aaw97414 KIAA0223	Aaw99198 DH cell K	Adh40334 Human min		99197 Min	Description		

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25
24	24	24	24	24	24	24	24	24	24	24	24	24	24	24	24	24	24	24	25	25
57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	59.5	
15	15	15	15	15	14	14	13	13	13	13	13	12	12	11	10	9	9	7	15	14
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ABR30612	ABR30473	ABR31271	ABR30940	ABR30509	ADR73051	ABG74854	ADD23767	ADG66270	ADG66271	ADG66272	ADG66269	ADD23610	AAW64577	ADD23269	ABR04849	ADM18218	ADM18256	AAB81550	ABB07889	ADU38989
Abr30612	Abr30473	Abr31271	Abr30940	Abr30509	Adr73051	Abg74854	Add23767	Adg66270	Adg66271	Adg66272	Adg66269	Add23610	Aaw64577	Add23269	Abr04849	Adm18218	Adm18256	Aab81550	Abb07889	Adu38989
_	Human can	Human can	Human can	Human can	L plantar	Human CHP	Breast ca	Human pro	Human pro	Human	Human pro	Breast ca	Human Fac	Breast ca	Human can	Chlamydia	C trachom	Zif268 zi	Phosphoen	Mutant S1

ALIGNMENTS

AAW99197 standard; peptide; 9 AA.

RESULT 1
AAW9197
ID AAW9
XX AAW9
AC AAW9
AC AAW9
AC Mino
XX USP
XX COPP
XX COP Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease. Goulmy EAJM, Hunt DF, 23-JUL-1997; 23-JUL-1998; 04-FEB-1999. WO9905174-A1 Minor histocompatibility antigen HA-1 T-cell epitope #3 20-MAY-1999 AAW99197; Homo sapiens. (UYLE-) RIJKSUNIV LEIDEN. (first entry) 97EP-00202303 98WO-NL000425. Engelhard VH;

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation. Disclosure; Page 15; 47pp; English. treatment of severe aplastic anaemia, leukaemia, HA-1 - useful to treat immune versus graft disease in bone and immune deficiency

WPI; 1999-153312/13.

Sequence 9 AA; diseases

DB 2;

Length 9;

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RESULT 2
AAW97374
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RESULT 3
ADH40334
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AC ADH4
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                                                                                                                                                                                                                The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW97374 standard;
                                                                                                                                                                                                                                                                                                                                                                                  Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. detection of genetic aberrances.
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02-JUN-1998;
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  ADH40334;
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                         ADH40334 standard; peptide; 9 AA.
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ilarity 100.0%;
Conservative 0
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Pred. No. 2e+06;
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RESULT 4
AAW99198
ID AAWS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUN-2002; 2002EP-00013423.
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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                                                                                                   20-MAY-1999
                                                                                                                                                                    AAW99198 standard; peptide; 13
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                                                                  KIAA0223 protein
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                                                                                                   (first entry)
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                                                                    sequence.
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Pred. No. 2e+
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          Goulmy EAJM,
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                                                                                                                                                                transplant rejection; Graf neoplastic haematopoietic
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                                                                          23-JUL-1998;
                                                                                                                      WO9905173-A1
                                                                                                                                                                          T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease;
                                                                                                                                                                                                                                                                                 AAW97414 standard; protein; 13 AA.
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                                                                                                                                                                                                                                      20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 13
                               (UYLE-) RIJKSUNIV LEIDEN.
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        Engelhard VH;
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Pred. No.
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RESULT 6
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The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                  Claim 3;
                                                                                                                                                                                                                                                                A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine;
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                                                                                                                                                                                                                                                                                            HA-1 - useful to treat immune versus graft disease in bone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HA-1 H-allele sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW97375 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9905313-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; 85.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-MAY-1999
                                                                                                                                                                                                                                                  The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. detection of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                          Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 VLRDDLLEA 9
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  VLHDDLLEA 9
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                                                                                                         Conservative
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98EP-00870125.
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Pred. No.
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2e+06;
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RESULT 9
AAW99199
ID AAWS
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AC AAWS
AC AAWS
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AC AAWS
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XX
                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel method for providing epitopes of allelic CC variants of antigenic proteins from specific species based on single CC nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein flentifying, CC selecting allelic peptide/protein variants, expression product or its CC fragment encoded by DNA sequence having SNP, creating variant epitopes, CC selecting epitopes binding to MHC protein. A protein of the invention has CC cytostatic activity, and may have a use in a vaccine. The method is CC given species by applying the method for several protein from the CC individuals, where the SNP profile of one or more individuals from the CC cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer in the superior profile. A method of the invention is useful for transplanting haematopoietic stem cells from a donor to a superior to a server of the progression, regression or onset of a treated disease. The present considerable is used in the exemplification of the invention.
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                                                                                                                                                                                                                                                        Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-JUN-2003; 2003WO-EP006251.
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                                                                                      standard;
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                                                                                      peptide;
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Pred. No. 2e+0
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2e+06;
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20-MAY-1999

(first entry)

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RESULT 10
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Best Local :
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 23-JUL-1997;
                       23-JUL-1998;
                                                04-FEB-1999.
                                                                        WO9905173-A1
                                                                                               Homo sapiens
                                                                                                                      T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis neoplastic haematopoietic cell; KIAA0223 polymorphism.
                                                                                                                                                                   KIAA0223 polymorphism in HA-1 positive homozygous individuals.
                                                                                                                                                                                                                                               AAW97415 standard;
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                     the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 31; 47pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           new minor histocompatibility antigen,
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3 VLHDDLLEA 11
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                                                                                                                                                                                                                                                                                                                                                                                                             AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                              (first entry)
97EP-00202303
                       98WO-NL000424.
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                                                                                                                                                                                                                                               protein; 13 AA.
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88.9%;
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                                                                                                                                                                                                                                                                                                                                                                      Score 37;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               versus graft disease in bone
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                                                                                                                                                                                                                                                                                                                                                                                 Length 13;
                                                                                                                                autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 11
AAW99195
ID AAW99
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents the KIAA0223 polymorphism in HA-1 positive homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HIA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
             A new minor histocompatibility
                                                                                                                                                                                                                                                                                                                    Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia: varcina.
                                                              Goulmy EAJM,
                                                                                                                23-JUL-1997;
                                                                                                                                                                   04-FEB-1999
                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                         23-JUL-1998;
                                                                                                                                                                                          WO9905174-A1
                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                          Minor histocompatibility antigen HA-1 T-cell epitope #1
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                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW99195 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   versus-host disease.
                                                                                       (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunogenic peptide from minor histocompatibility antigen HA-1 for inducing tolerance to transplants and prevent rejection or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-142855/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Goulmy EAJM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VLRDDLLEA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.1%;
Similarity 88.9%;
8; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VLHDDLLEA 11
                                                                                                                                                                                                                                                                                                          aplastic
                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                              Hunt DF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hunt DF,
                                                                                                                97EP-00202303
                                                                                                                                         98WO-NL000425.
                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                       disease; bone marrow transplant; leukaemia; vaccine;
ic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       57pp;
                                                                                                                                                                                                                  His, Arg
                                                              Engelhard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Engelhard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 37; DB
Pred. No. 1.7;
antigen, HA-1 - useful to treat immune and host versus graft disease in bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                              ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e graft-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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and prevent rejection

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RESULT 12
AAW97572
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the particular it can be used with bone marrow transplantation, in the particular it can be used with bone marrow transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            marrow
                          The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HIA class I context, can be eliminated after specific recognition of the peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T-cell epitope; minor transplant rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW97572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW97572 standard; peptide;
                                                                                                                                                                                                                                                                        Goulmy
                                                                                                                                                                                                                                                                                                                                    23-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                 23-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                               04-FEB-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                             WO9905173-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neoplastic haematopoietic cell.
                                                                                                                                                                             Immunogenic peptide fro for inducing tolerance versus-host disease.
                                                                                                                                                                                                                                             WPI; 1999-142855/12
                                                                                                                                                   Claim 1; Page 39; 57pp;
   peptides can also be used to raise T-cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       epitope from the minor histocompatibility antigen HA-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
8; Conserv
                                                                                                                                                                                                                                                                          EAJM,
                                                                                                                                                                                                                                                                                                       RIJKSUNIV LEIDEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                        Hunt DF,
                                                                                                                                                                                                                                                                                                                                     97EP-00202303
                                                                                                                                                                                                                                                                                                                                                                   98WO-NL000424.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              minor histocompatibility antigen HA-1; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "His or Arg"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 47pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               aplastic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85.7%;
88.9%;
                                                                                                                                                                                              from minor histocompatibility antigen HA-1 - nce to transplants and prevent rejection or gr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Graft-versus-Host Disease;
                                                                                                                                                    English
                                                                                                                                                                                                                                                                          Engelhard
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               th bone marrow transplantation, in the anaemia, leukaemia, and immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                    antibodies,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                    T-cell
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AD773023
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                                                                          CC encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC expressition comprising an above peptide mixed with a pharmaceutical CC expression vector with a nucleic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC Tymphocytes (CTLs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (prophylactic or therapeutic) vaccine composition for a DNA-based CC immunisation, or for preparing an immune response provoking vaccine in CC the event of RSV infection (the vaccine being prepared by contacting the CC polypeptide in an immune response-provoking and activity. This cappide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADT73023 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human respiratory syncytial virus; cytotoxic T lymphocyte; MHC epitope; DNA-based immunisation; RSV infection; virucio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-JAN-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-2004; 2004WO-EP004061.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-OCT-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-758334/74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-2003; 2003EP-00447095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to novel isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 16; SEQ ID NO 919; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lasters I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ALGO-)
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8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALGONOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Desmet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                high affinity binding peptide SeqID 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       syncytial virus
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RSV),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RSV infection; virucidal; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or purified peptides of the in particular ten RSV genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lar ten RSV genes
proteins NS-1 (also
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Query Match
Best Local Similarity
Matches 4; Conserv

Conservative

4 Score Pred.

69.0%;

core 29; DB 8; red. No. 2e+06; Mismatches

Length

9

Indels

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Gaps

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Sequence 9

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RESULT 14
ANAW47427
ID ANAW47
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XX II-DI
XX II-DI
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XX III-DI
RESULT 15
ADT73690
ID ADT73
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AC ADT73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 4
                      Human respiratory syncytial virus
                                                                           human respiratory syncytial virus; cytotoxic T lymphocyte; CTL; MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
                                                                                                                                                                                                                                                                       ADT73690;
                                                                                                                                                    Human RSV L high affinity binding peptide SeqID 1586.
                                                                                                                                                                                                               13-JAN-2005
                                                                                                                                                                                                                                                                                                                    ADT73690 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence from preserved region VI of prenyl diphosphate synthetase (PDS), was used in the preparation of primers for the amplification of PDS DNA. Substances synthesized by PDS, i.e. polyprenyl diphosphates, are precursors of physiologically active substances, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vitamin K and ubiquinones
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This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC composition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC Tlymphocytes (CTLs) that respond to a major histocompatibility complex (CMHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC RSV peptide is useful for preparing a diagnostic composition or an RSV immunisation, or for preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the communisation for a fire-preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the
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                                                                           Sequence 9 AA;
                                                                                                                               polypeptide in an immune response-provoking amount of specific CTL). Accordingly, these peptide compositions have virucidal activity. This peptide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 16; SEQ ID NO 1586; 143pp; English.
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	re 19.5; DB 2; d. No. 1.2e+03; Mismatches 1;	on 13-Mar-1997 #text_change M.; Kuboki, Y. 1996 rells synthesized C-terminal 86332; PMID:9244181 UNIPARC:UPI000017C63D		Score 21; DB 2; L Pred. No. 6.2e+02; 2; Mismatches 1;	(216347; PMID:8163531 UNIPARC:UPI000017C62E	d alpha-6 asso)ec-1995	ALIGNMENTS	2876	0 Q 4 A		.
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R;Fubhitani, K.; Higashiyama, K.; Asao, M.; Hosokawa, K.
Biochim. Biophys. Acta 1292, 273-280, 1996

A;Title: Characterization of the constituent polypeptides of A;Reference number: S65721; MUID:96176855; PMID:8597573

A;Accession: S65730
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C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Graves, P.E.; Kaminsky, L.S.; Halpert, J.
Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional and structural multiplicity
A;Reference number: A26997; MUID:88000604; PMID:3651420
A;Accession: C26997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unspecific monooxygenase (EC 1.14.14.1) isozyme E, phenobarbital-inducible, hepatic - N;Alternate names: cytochrome P450 C;Species: Rattus norvegicus (Norway rat) C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 31-Dec-2004 C;Accession: C26297
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A;Title: Characterization of genes that encode subunits of cucumber A;Reference number: A56819; MUID:91355209; PMID:1883835
A;Accession: E56819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PS I complex subunit 8 - cucumber (fragment)
C;Species: Cucumis sativus (cucumber)
C;Date: 05-Jan-1996 #sequence_revision 05-Jan-1996 #text_change 09-Jul-2004
     Ś
                                                                                                                                                                                                                                                                                        hemoglobin, extracellular, component - earthworm (Lumbricus terrestris) (fr C;Species: Lumbricus terrestris (common earthworm) (C;Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
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                                                                                                                A; Cross-references: UNIPARC: UPI000017BD88
                                                                                                                               A; Molecule type: protein A; Residues: 1-12 < FUS>
                                                                                                                                                                 A; Status: preliminary
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Pred. No.
                                         2; Mismatches
                                                          Score 18;
Pred. No.
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PH0924
T-cell receptor beta chain V-D-J region (isolate 10) - rat C; Species: Rattus norvegicus (Norway rat)
C; Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_ch
C; Dacession: PH0924
C; Accession: PH0924
L; Sun. D.; Wiley, S.; Vandenbark, A
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A;Title: Purification and characterization of the proteinase ECP 32 from Es. A;Reference number: S71920; MUID:96350420; PMID:8765229
A;Accession: S71920
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A;Residues: 1-15 <GRA>
A;Residues: UNIPARC:UPI0000174DA7
A;Cross-references: UNIPROT:Q06884; UNIPARC:UPI0000174DA7
C;Superfamily: Cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N;Alternate names: cytochrome P450b
c;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Jun-1988 #sequence_revision 30
C;Accession: A26997
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A;Experimental source: strain A2
C;Keywords: hydrolase; metalloproteinase
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C;Species: Escherichia coli
C;Species: Escherichia coli
C;Date: 04-Feb-1998 #sequence_revision 13-Feb-1998 #text_change 09-Jul-2004
C;Accession: S71920
A;Cross-references: UNIPARC:UPI000017C9F3
A;Experimental source: concanavalin A-activated lymphoblast C;Keywords: T-cell receptor
                                                                                                                             A;Accession: PH0924
A;Molecule type: mRNA
A;Residues: 1-11 <GOL>
                                                                                                                                                                                                                                                                                                  R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta
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A; Accession: A26997
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Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional
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                                                                                                                                                                                                                                                      A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental A;Reference number: PH0891; MUID:92078857; PMID:1836012
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1; Mismatches
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 19-May-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: S39762
R;Ohishi, N.; Imaoka, S.; Suzuki, T.; Funae, Y.
Biochim. Biophys. Acta 1158, 227-236, 1993
A;Title: Characterization of two P-450 isozymes placed in the rat CYP2D sub A;Reference number: S39761; MUID:94072607; PMID:8251521
A;Accession: S39762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 10
S39762
A;Status: nucleic acid sequence A;Molecule type: mRNA A;Residues: 1-13 <ADE>
                                                     R;Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R. Neuron 9, 209-216, 1992
A;Title: Calcium-activated potassium channels expressed A;Reference number: JH0697; MUID:92360298; PMID:1497890
A;Accession: PS0443
                                                                                                                                                 potassium channel protein Slo G3 - fruit fly (Drosophila melanogaster) (fragcises: Drosophila melanogaster) (Fragcises: Drosophila melanogaster) (Fragcises: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Feb-1997 (Fragcession: PS0443)
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Cell 62, 591-598, 1990
A;Title: The c4 repressors of bacteriophages P1 and P7 are
A;Reference number: $42448, MUID:90335968; PMID:1696181
A;Accession: $42449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: phage P7
C;Date: 07-Sep-1994 #sequence_revision 26-May-1995 #text_change 09-Jul-2004
C;Accession: $42449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytochrome P450 UT-7b - rat (fragment)
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Matches 3
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;Residues: 1-11 <CIT>
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                                                                                                                             K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta,
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Pred. No. 3e+03;
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Pred. No.
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Pred. No. 2.7e+03;
1; Mismatches 0
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S57584
T cell receptor
C;Species: Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chlorophyll a/b-binding protein Lhcb5 - spinach (fragment)
N;Alternate names: light-harvesting complex LHCIIc protein
C;Species: Spinacia oleracea (spinach)
C;Date: 19-Mar-1997 #sequence_revision 23-Apr-1999 #text_change 23-Apr-1999
C;Accession: S50900
R;Walters, R.G.; Ruban, A.V.; Horton, P.
Bur. J. Biochem. 226, 1063-1069, 1994
A;Title: Higher plant light-harvesting complexes LHCIIa and LHCIIc are bound A;Reference number: S50900, MUID:95112835; PMID:7813461
A;Accession: S50900
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S50900
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C;Keywords: oxidoreductase
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A; Accession: C39170
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C;Superfamily: chlorophyll a/b-binding protein
C;Keywords: chlorophyll; chloroplast; light-harvesting complex;
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C;Comment: This potassium channel is activated by calcium.
C;Genetics:
C;Genetics:
A;Gene: FlyBase:slo
A;Cross-references: FlyBase:FBgn0003429
C;Keywords: alternative splicing; ion channel; potassium channel; transmembrane
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100.0%; Pred. No. 3.5e+03;
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C;Date: 19-Oct-1995 #sequence_revision 17-Nov-1995 #text_change 05-Nov-1999
C;Accession: 857584
R;Burrows, S.R.; Silins, S.L.; Moss, D.J.; Khanna, R.; Misko, I.S.; Argaet, V.P.
submitted to the EMBL Data Library, June 1995
A;Description: T cell receptor repertoire for a viral epitope in humans is diversified A;Reference number: 857494
A;Recession: 857584
A;Status: preliminary
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-15 *SURA
A;Cross-references: UNIPARC:UPI0000116747; EMBL:Z49956; NID:g887466; PIDN:CAA90227.1; PC;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hucolin, 75K chain - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
C;Accession: S68004
R;Edgar, P.F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 15
S68004
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A;Title: Hucolin, a new corticosteroid-binding protein from human plasma with structural A;Reference number: S68004; MUID:96087107; PMID:7498469
A;Accession: S68004
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 < EDG>
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Search completed: April Job time : 20 secs
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                                                                                                                                                                                     Query Match 38.1%; Score 16; DB 2; L
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 3; Conservative 0; Mismatches 0;
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4 DDL 6
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Maximum Match 100%
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Maximum DB seq
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Gapop 10.0 , Gapext 0.5
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      2166443 seqs, 705528306 residues
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Copyright (c) 1993 - 2006 Biocceleration
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Q47600 ECOLI
Q718N9 PARA
UPO5 MÖUSE
Q718K6 PARA
UPC7 MAIZE
Q71063 MOUSE
P82560 STRPY
Q7M063 MOUSE
P82560 STRPY
Q7M062 MOUSE
Q9UMM5 SULSO
Q9UMIS HUMAN
Q9UC60 CICOS
Q9R556 CHVI
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RESULT 2 Q718N9 9PARA ID Q718N9; PRT; 10 AA. AC Q718N9; DT 05-UUL-2004 (TrEMBLrel. 27, Created) DT 05-UUL-2004 (TrEMBLrel. 27, Last sequence update) DT 05-UUL-2004 (TrEMBLrel. 27, Last annotation updat DE Hemagglutinin-neuraminidase (Fragment). OS Newcastle disease virus. OC Viruses; ssRNA negative-strand viruses; Mononegav Paramyxoviridae; Paramyxovirinae; Avulavirus. OX NCBI_TaxID=11176; RN NUCLEOTIDE SEQUENCE. RA Gould A.R., Hansson E., Selleck K., Kattenbelt J. RA Della-Porta A.J.; RI "Newcastle disease virus fusion and hemagglutinin RT motifs as markers for viral lineage."; RL Avian pathol. 32:361-373(2003). DR EMBL, AF542884; AAQ11609.1; -; Genomic_DNA.	Query Match 59.5%; Score 25; DB 2; Le Best Local Similarity 62.5%; Pred. No. 5.5e+02; Matches 5; Conservative 1; Mismatches 2; Qy 2 LRDDLLEA 9	RESULT 1 Q47600_BCOLI Q47600_BCOLI PRELIMINARY; PRT; 11 AA. AC Q47600; DT 01-NOV-1996 (TrEMBLrel. 01, Created) DT 01-NOV-1996 (TrEMBLrel. 19, Last sequence update) DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation updat DE REase; GN Name=ERase; GN Name=FRAse; CS Escherichia coli. OC Escteria; Proteobacteria; Gammaproteobacteria; En OC MICHEOTICE SEQUENCE. RN NUCLEOTIDE SEQUENCE. RN MEDLINE=91139577; PubMed=1995588; RN MEDLINE=91139577; PubMed=1995588; RT Tao T.; Bourne J.C., Blumenthal R.M.; RT Tao T.; Bourne J.C., Blumenthal	32 18 42.9 13 2 QTTE26 9CLOS 33 18 42.9 13 2 QTTE30—9CLOS 34 18 42.9 13 2 Q957T7—9PERC 35 18 42.9 14 2 Q758BC MAIZE 36 18 42.9 14 2 Q758BC ECOLI 37 18 42.9 15 2 Q7XBBC ECOLI 39 17 40.5 10 2 Q55732—9FUNG 40 17 40.5 10 2 Q55732—9FUNG 41 17 40.5 11 2 Q51K70—HUNAN 43 17 40.5 11 2 Q51K70—HUNAN 43 17 40.5 11 2 Q51K70—HUNAN 44 17 40.5 12 2 Q7GCS7—ORYSA 45 17 40.5 12 2 Q7GCS7—ORYSA 46 17 40.5 12 2 Q7GCS7—ORYSA
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01-OCT-1994 (Rel. 30, Created)
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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                                                                                  Della-Porta A.J.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct protein sequencing.
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"Newcastle disease virus fusion and he motiff as markers for viral lineage."; Avian Pathol. 32:361-373(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                               Q718K6 PPARA PRELIMINARY;
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                                                                                                                                                                           NCBI_TaxID=11176;
                                                                                                                                                                                                                          Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Avulavirus.
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Newcastle disease virus.
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                                                             hemagglutinin-neuraminidase gene
                                                                                                                  Kattenbelt J.A., Mackenzie M.,
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RESULT 6
Q7M063 MOUSE
Q7M063 MOUSE PRELIMINARY;
AC Q7M063;
AC Q7M063;
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Calnexin (Fragment).
OS Mus musculus (Mouse).
OC Eukaryota, Metazoa; Chordat
OC Muridae; Murinae; Mus.
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01-OCT-1996 (Rel.
25-OCT-2004 (Rel.
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Pernollet J.-C., Zivy M., de V.
"The maize two dimensional gel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Trach
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                      Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANEOUS: On the 2D-gel the dete protein is: 6.4, its MW is: 48.4 kDa.
                                                                                                                                                                                                                                                                                                                                                                                                                     genome analysis program.";
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             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
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RESULT 8
Q7M062 MOUSE
Q7M062 MOUSE PRELIMINARY;
AC Q7M062;
DT 01-MAR-2004 (TYEMBLrel. 26,
DT 01-MAR-2004 (TYEMBLrel. 26,
DT 01-MAR-2004 (TYEMBLrel. 26,
DT 01-MAR-2004 (TYEMBLrel. 26,
DE COllagen type I (Fragment).
OS Mus musculus (Mouse).
OC Musmaryota; Metazoa; Chordata
OC Mammalia; Sutheria; Euarchor
OC Muridae; Murinae; Mus.
OX NCBI_TAXID=10090;
RN [1]
RP PROTEIN SEQUENCE.
RX MEDLINE=97386332; PubMed=92.
RA Mizuno M., Kitafima T., Tom:
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P82560;
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01-MAR-2004 (TrEMBLrel 26,
01-MAR-2004 (TrEMBLrel 26,
Collagen type I (Fragment).
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                                                                                                                                   Collagen type I (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Euarchonto
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STRAIN=JRS4;
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Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
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01-OCT-2003 (TrEMBLrel. 25, I
Unknown protein from 2D-page
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MEDLINE=94216347; PubMed=8163531;
Lenter M., Vestweber D.;
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J. Biol. Chem. 269:12263-12268(1994).
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VanBogelen R.A.;
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    PubMed=9244181; DOI=10.1016/0167-4889(95)00151-4; ma T., Tomita M., Kuboki Y.;
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37.5%;
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-page (Fragment).
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RESULT 9
QSUMM5 SULSO
AC QSUMM5;
DT 01-MAY-2000 (
DT 01-MAY-2000 (
DT 01-JUN-2022 (
DE Ribosomal pro
OS Sulfolobus so
OC Archaea; Cren
OC Archaea; Cren
OC Archaea; Cren
CR [1] TaxID=22
RN [1] FROTEIN SEQUE
RY [1] FROTEIN SEQUE
RY [1] FROTEIN CO.; TaxID=22
RN [1] FROTEIN CO.;
RESULT 10
QUUNLB | HUMAN
ID QUUNLB |
AC QUUNLB |
DT 01-MAY-2
DT 01-MAY-2
DT 01-MAY-2
DT 01-MAY-2
DT 01-MAY-2
OG Name=APC
GS HOMO SAP
OC ENKARYOL
OC MAMMAIA
OC HOMO .
OX NCBI_Tax
RN [1]
RA Carr I.
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Matches 3
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Best Local S
Matches 6
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QSUNLB;

QSUNLB;

O1-MAY-2000 (TrEMBLrel. 13, C

O1-MAY-2000 (TrEMBLrel. 13, L

O1-MAY-2000 (TrEMBLrel. 13, L

APC2 protein (Fragment).
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                                             Submitted (DEC-1998) to the EMBL; AF110338; AAD29275.1; NON_TER 1 1
                                                                                                       NUCLEOTIDE SEQUENCE.
Carr I.M., Markham A.F.,
Meredith D.M.;
                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                          Name=APC2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neighbors.";
J. Biol. Chem. 266:21578-21583(1991).
NON_TER 10 10
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "The osteoblastic MC3T3-E1 cells synthesized C-t type I collagen, which promoted cell-attachment Biochim. Biophys. Acta 1310:97-102(1996).
PIR: S62675; S62675.
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                                                                                                                                                                                                                                                                        Homo sapiens (Human)
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15 AA; 1729 MW;
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L12e, Sso L1(
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Q47602;
Q47602;
Q1-NOV-1996 (TrEMBLrel. 01,
01-NOV-1996 (TrEMBLrel. 01,
01-DEC-2001 (TrEMBLrel. 19,
REase protein (Fragment).
Name=REase;
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09ZB42;
01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "A family of regulatory genes associated with type modification systems.";
J. Bacteriol. 173:1367-1375(1991).
EMBL; M63620; AAA24558.1; -; Genomic_DNA.
NON_TER 11 11
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MEDLINE=91139577; PubMed=1995588;
Tao T., Bourne J.C., Blumenthal R.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria; Gammaproteobacteria;
Enterobacteriaceae; Escherichia.
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NCBI_TaxID=1314;
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                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                    streptococci.
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"Characterization of nra, a ;
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Bacteria; Firmicutes; I
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                                                                                             Local Similarity
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1. Microbiol. 31:0-0(1999).
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Pred.
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Pred. No. 1.1e+04;
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J. 8.5e+03;
J. 0;
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P42052;
01-NOV-1995
01-NOV-1995
05-JUL-2004
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Photosystem I; T
NON_TER 15
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; agnoliophyta; eudicotyledons; core eudicotyledons; rosids; eurosids \mathbf{I}_i Cucurbitales; Cucurbitaceae; Cucumis.
                      PROTEIN SEQUENCE.

MEDILINE=95392215; PubMed=7663175; DOI=10.1006/prep.1995.1049;
De Pauw P., Neyt C., Vanderwinkel E., Wattiez R., Falmagne P.;
De Pauw P., Neyt C., Vanderwinkel E., Wattiez R., Falmagne P.;
"Characterization of human serum N-acetylmuramyl-L-alanine ami
purified by affinity chromatography.";
Protein Expr. Purif. 6:371-378(1995).
GO; GO:0008745; P:N-acetylmuramyl-L-alanine amidase activity.
GO; GO:0001519; P:peptide amidation; NAS.
SEQUENCE 15 AA; 1600 MW; 9016B00PFF9E780A CRC64;
                                                                                                                                                                                                                              01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
N-acetylmuramyl-L-alanine amidase (EC 3.5.1.28) (Fragment).
                                                                                                                                                                                                                                                                                     Q9UC60 HUMAN PRELIMINARY;
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Chloroplast; Direct protein
Photosystem I; Thylakoid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restructed the Entropean Bioinformatics Institute.
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"Characterization of genes that encode subunits of cucumber PS I complex by N-terminal sequencing.";
Biochim. Biophys. Acta 1059:141-148(1991).
-i- FUNCTION: Essential for the activity of NADP photoreduction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=3659;
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                                                                                                                                                                                                       Eukaryota; Metazoa;
                                                                                                                                                                                                                  Homo sapiens
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RESULT 15
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O1-MAY-2000 (TrEMBLrel. 13, Created)
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O1-MAY-2000 (TrEMBLrel. 14, Last amotation update)
DT
O1-JUN-2000 (TrEMBLrel. 14, Last amotation update)
DE
BCL2-interactive cell death susceptibility regulator (Fragment).
OS
ENCATYOTIS; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
OC
MCBI_TaxID=9606;
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MCBI_TaxID=9606;
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McJerink J.P., Smetsers T.F., Sloetjes A.W., Linders E.H.,
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Mensink E.J.;
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Mensink E.J.;
RA
Mensink E.J.;
RT
malignancies,;
RL
Leukemia 9:1028-1832(1995).
SQ
SEQUENCE
15 AA; 1860 MW; 4F9238D6FE9743DB CRC64;
Search completed: April 6, 2006, 17:02:57 Job time : 116.333 secs
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                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
NUCLEOTIDE SEQUENCE.
NUCLEOTIDE SEQUENCE.
NEDLINE=96071131; PubMed=7475270;
Meljerink J.P., Smetsers T.F., Sloetjes A.W., Linders E.H.,
Mensink B.J.;
Mensink B.J.;
"Bax mutations in cell lines derived from hematological
malignancies.";
Leukemia 9:1826-1832(1995).
SEQUENCE 15 AA; 1860 MW; 4F9238D6FE9743DB CRC64;
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4 IRDBL 8
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/iaa/H_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
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(c) 1993 - 2006 Biocceleration Ltd
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US-09-27-609A-11
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                                                                                                                                                                   Sequence 5, Application US/09489760
Patent No. 6678375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 198-07-23
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                              NUMBER OF SEQ ID NOS: 16
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LENGTH: 9
TYPE: PRT
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Best Local
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ORGANISM: KIAA0223 partial complementary DNA
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	Sequence 20.	Sequence 20,	Sequence 20,		Sequence 20,	Sequence 104	Sequence 79,	Sequence 104	Sequence 104	Sequence 104	Sequence 104	Sequence 4,	Sequence 104				
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ALIGNMENTS

; OTHER INFORMATION: Description of Artificial Sequence: US-09-269-250E-18 APPLICANT: Goulmy, Elsa TITLE OF INVESTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN FILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: US/09/269,250E CURRENT FILING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 18 Sequence 18, Application US/09269250E Patent No. 6830883 ORGANISM: Artificial Sequence FEATURE: 1 VLRDDLLEA 9 |||||||| 1 VLRDDLLEA 9 h 100.0%; Score 42; DB 2; Similarity 100.0%; Pred. No. 4.6e+05; 9; Conservative 0; Mismatches 0; Length 9; Indels Exon Fragments ٥, Gaps 0 HA-1

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US-09-269-250E-26
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
APPLICANT: Hard, Victor H
ITILE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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                                                                                                            ; OTHER INFORMATION: Description of Unknown Organism:HA-1-/-US-09-489-760-14
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SEQ ID NO 26
LENGTH: 13
TYPE: PRT
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APPLICANT: Goulmy, Elsa
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CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
RUMBER OF SEQ ID NOS: 38
RUMBER OF SEQ ID NOS: 38
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                                                                                                                                                                                     SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 14
LENGTH: 13
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                                                                                   Query Match
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                                                                                                                                       TYPE: PRT
ORGANISM: Unknown Organism
FEATURE:
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Similarity 100.0%;
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milarity 100.0%;
Conservative 0;
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Pred. No.
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                                                             Score 42; DB 2;
Pred. No. 0.028;
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GENERAL INFORMATION:

APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Hunt, Donald F

APPLICANT: Hunt, Donald F

APPLICANT: Hard, Victor H

TITLE OF INVENTION: The HA-1 Antigen
TITLE OF INVENTION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21

EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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US-09-269-250E-20
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CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOPTWARE: Patentin version 3.1
SEQ ID NO 20
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Patent No. 6830883
GENERAL INFORMATION:
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Best Local S
Matches 8
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
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GENERAL INFORMATION:

APPLICANT: GOULT, B1sa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
TITLE REFERENCE: 58994
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
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                                                                                                                Sequence 28, Application US/09269250E Patent No. 6830883
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0; Mismatches
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 16
LENGTH: 13
; ORGANISM: Human
; PEATURE;
; NAME/KEY: MISC_FEATURE
; LOCATION: (3)..(3)
; OTHER INFORMATION: Xaa represents a
US-09-269-250E-29
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US-09-489-760-16
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Matches 8; Conserve
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                                                                                                                 APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
LENGTH: 9
                                                                                                                                                                                                                                                                     Patent No. 6830883
GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Artificial Sequence
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GENERAL INFORMATION:
APPLICANT: Rajksuniversiteit Te Leidin
APPLICANT: Roulmy, Elsa A.J.M
APPLICANT: Hunt, Donald P.
APPLICANT: Hunt, Donald P.
APPLICANT: Hard, Victor H.
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT FILING DATE: 2000-01-21
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: DCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN Ver. 2.1
SOFTWARE: PATENTIN Ver. 2.1
SOFTWARE: PATENTIN Ver. 2.1
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US-09-217-609A-11
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Best Local Similarity
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APPLICANT: MURAMA
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APPLICANT:
APPLICANT:
        COUNTRY: US
ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windows
                                                                                                                                                                                               APPLICANT: KOYAMA, Tanetoshi
APPLICANT: SHIMIZU, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 9
CURRENT APPLICATION DATA
                                                                                                                                     STATE:
                                                                                                                                                   CITY: Washington
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1025 Connecticut Avenue,
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OGURA, Kyozo
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05
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US-08-873-2358-11
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                                                                                                                                                                                                                                                       ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,235B
FILING DATE: 11-Jun-1997
                                                                    APPLICATION NUMBER: JP 154441/1996
ETILING DATE: 14-Uun-1996
ATTORNEY/AGENT INFORMATION:
NAME: TOFFENETTI, Judith L.
REGISTRATION NUMBER: 39,048
REFERENCE/DOCKET NUMBER: 10235/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-429-1776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: MURAMA
APPLICANT: KOIKE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 202-429-0796
INFORMATION FOR SEQ ID NO:
           TELEFAX: 202-429-0796
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: KOITE, Ayumi
APPLICANT: OGURA, Kyozo
APPLICANT: KOYAMA, Tanetoshi
APPLICANT: SHIMIZU, Naoto
APPLICANT: GHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
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REFERENCE/DOCKET NUMBER: 10235/1
TELECOMUNICATION INFORMATION:
TELEPHONE: 202-429-1776
TELEPAX: 202-429-0796
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ATTORNEY/AGENT INFORMATION:
NAME: TOFFENETTI, Judith L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
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APPLICATION NUMBER:
                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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TOPOLOGY: linear
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1025 Connecticut Avenue, NW -
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Pred. No. 4.6e+05;
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Best Local Similarity
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US-09-269-250E-38
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                                                                                                US-09-972-115A-64
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CURRENT PILING DATE: 199-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 38
LENGTH: 8
LENGTH: 8
                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/129,123
PRIOR FILING DATE: 1999-04-13
NUMBER OF SEQ ID NOS: 64
SOFTWARE: Patentin version 3.1
SEQ ID NO 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Patent No. 6830883
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                                   Matches
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                                                             Query Match
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/972,115A
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/128,577
PRIOR FILING DATE: 2000-04-10
                                                                                                                                                                                                                                                                                                                                         APPLICANT: Geron Corporation
APPLICANT: Gregg, Morin B.
APPLICANT: Walter, Funk D.
APPLICANT: Maler, Funk D.
                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: A Second Mammalian Telomerase FILE REFERENCE: 080/003C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
                                                                                                              LENGTH: 11
TYPE: PRT
ORGANISM: Homo sapiens
                                                  Local Similarity
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100.0%; Pred. No. 4.6e+05;

 Mismatches

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                                                  Score 23; DB 2;
Pred. No. 1.7e+02;
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Pred. No. 4.6e+05;
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US-08-962-284-8

US-08-962-284-8

Sequence 8, Apsilcation US/08962284

Patent No. 598568

Patent No. 598568

GENERAL INFORMATION:

APPLICANT: Land, Bliaabeth J.

APPLICANT: Pestconjamasp, Kersi N.

APPLICANT: Mulffulle, Julia D.

TITLE OF INVERTION: AND NUCLEIC ACIDS ENCODING THE SAME

UNDERSORES FISHS & KINATAGON P.C.

STREET: 225 Franklin Street

COMPUTER: READABLE PORN:

MEDIUM TYPE; Diskette

COMPUTER: APPLICATION DATA:

APPLICATION UNDERS: 32.093

REFERENCE/DOCKET NUMBER: 07917/058001

TELEPAN: 617/542-5070

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Minimum DB
Maximum DB
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Maximum Match 100%
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Perfect score:
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 0.5
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

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4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

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US-11-077-740-18
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US-10-623-176-74
US-10-623-176-74
US-10-623-176-74
US-10-623-176-7
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                 Sequence
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  Sequence
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18, Appli
18, Appli
18, Appli
14, Appli
26, Appli
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46, Appli
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RESU US-1	p 8	Qu Be	<u> </u>	SEQ SEQ TI	יייייייייי	יי יי יי יי יי		RESU US-1 ; Pu					
RESULT 2 US-10-791-2:		Query Match Best Local Matches	FEATURE: OTHER INFO FEATURE: NAME/KEY: LOCATION: 10-623-176:	ID ID SNGT	PRIOR FI PRIOR AP PRIOR AP PRIOR FI	URRENT RIOR AP	SENERAL INFO APPLICANT: APPLICANT: APPLICANT: TITLE OF IN	RESULT 1 US-10-623-176-10 ; Sequence 10, Applicat; Publication No. US300;		441 443 444 45	40 40	7 C T T C	29 31 31
5, Ap	A = A	5	: NFORMAT: Y: SITE Y: (1). 76-10	SEC Pat O Pat O E A	FILING APPLICA FILING APPLICA APPLICA FILING	APPLI FILIN FILIN PLICA	T: GO	76-10 10, A		22222	N N N N N	2288	29 29 29
5 Application	VLRDDLLEA 9 VLRDDLLEA 9	Similarity 9; Conservat	ION:	NOS: In Ver	FILING DATE: 1997-07-2. APPLICATION NUMBER: PC FILING DATE: 1998-07-2. APPLICATION NUMBER: JP APPLICATION NUMBER: JP FILING DATE: 2000-01-2.	CURRENT APPLICATION NUMBER: US/1 CURRENT FILING DATE: 2003-07-18 PRIOR APPLICATION NUMBER: 09/489 PRIOR FILING DATE: 2000-01-21 PRIOR FILING DATE: 2000-01-21	Goulmy, Els A Hunt, Donald Hunt, Donald Engelhard, V IVENTION: HA-1	.0 Application US/10623176 0. US20040092446A1		59.5 57.1 57.1 57.1 57.1	9.9.	- 4 6 6 6	
g		100. 100. tive	Description	77 . 2.1 Sequenc		NUMBER: US/ 3: 2003-07-1 3: 2003-07-1 1UMBER: 09/48 2000-01-21	A.J.M. ld F. Victor	m US,		10 7 9	10	11111	13 9 14
107		8 %	òt i	ä	7-23 PCT PCT 1-23	-21 -21	or M	/10 46A		04400			
/10791217		<pre>\$; Score 42; DB 4; Length b; Pred. No. 1.7e+06; 0; Mismatches 0; Indel</pre>	of Artificial Sequence:	ĕ	-24 PCT/NL98/00424 -23 JP 2000-504165	0/623, ,760	c H. copes and uses thereof	0623176 11	ALIGNMENTS	US-10-805-650-90 US-10-271-708-16 US-10-623-176-4 US-10-503-135-26 US-10-503-135-64	US-10-153-344-111 US-09-834-765-349 US-10-623-176-39 US-10-702-400-90	US-10-623-176-40 US-10-623-176-6 US-10-623-176-54 US-10-623-176-8 US-10-623-176-8	US-10-623-176-64 US-10-623-176-5 US-10-623-176-47 US-10-623-176-62
		19; els 0; Gaps	HA-1 peptide							16 16 26 64	Sequence 11, Appl Sequence 349, Appl Sequence 39, Appl Sequence 90, Appl	Sequence 40, Appli Sequence 54, Appli Sequence 54, Appli Sequence 8, Appli	Sequence 64, Appl Sequence 5, Appli Sequence 47, Appl Sequence 62, Appl
		0,											

Sequence 5, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H

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Sequence 44, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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; TYPE: PRT
; ORGANISM: Artificial sequence
; FEATURE:
; FEATURE:
; OTHER INFORMATION: obtained from KIAA0223 partial complementary
US-10-791-217-5
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US-11-007-740-18
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                                                                                                                                                                                                                                                                                            RESULT 4
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Best Local (
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CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: The HA-1 Antigen FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 40
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PRIOR APPLICATION NUMBER: 09/489,760 PRIOR FILING DATE: 2000-01-21 PRIOR APPLICATION NUMBER: EP 97202303.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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Pred. No. 1.7e+06;
; Mismatches 0;
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GENERAL INFORMATION:

APPLICANT: Goulmy, Els A.J.M.

APPLICANT: Goulmy, Els A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

TITLE OF INVENTION: HA-1 epitopes and uses thereof

FILE REFERENCE: 2183-6047US

CURRENT PILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2000-01-21

PRIOR APPLICATION NUMBER: B97202303.0

PRIOR APPLICATION NUMBER: B97202303.0

PRIOR APPLICATION NUMBER: B97202303.0

PRIOR APPLICATION NUMBER: D97202303.0

PRIOR APPLICATION NUMBER: D7723

PRIOR APPLICATION NUMBER: D770128

PRI
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US-10-623-176-65
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SEQ ID NO 65
LENGTH: 12
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SEQ ID NO 44
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PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
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NAME/KEY: SITE
LOCATION: (1)..
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NUMBER OF SEQ ID NOS: 77
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                                                                                              NAME/KEY: SITE LOCATION: (1).
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TYPE: PRT
ORGANISM: Artificial Sequence
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OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                              Local Similarity
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                                  VLRDDLLEA 9
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VLRDDLLEA
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Pred. No. 0.26;
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Pred. No. 0.31;
Mismatches
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US-10-623-176-74 ; Sequence 74, Application US/10623176 ; Publication No. US20040092446A1

RESULT 6

GENERAL INFORMATION:

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FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: DCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER: OF SEO ID NOS: 17
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 14
LENGTH: 13
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                                                                                                                                                                           ; OTHER INFORMATION: obtained from HA-1-/- phenotype US-10-791-217-14
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Best Local Similarity
Matches 9; Conserve
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US-10-623-176-74
                                                                              Query Match 100.0%; Score 42; DB 4; Length 13; Best Local Similarity 100.0%; Pred. No. 0.34; Matches 9; Conservative 0; Mismatches 0; Indels
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APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OP INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Domald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
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PRIOR TILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
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                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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ORGANISM: Artificial Sequence
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VLRDDLLEA 11
                           VLRDDLLEA 9
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100.0%; Pred. No. 0.34;
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-42
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Sequence 26, Application US/11007740
Sequence 26, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: ECT/NL98/00424
PRIOR APPLICATION NUMBER: 972000-504165
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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; SEQ ID NO 42
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CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.1
SEQ ID NO 26
LENGTH: 13
                                                                              Best Loc
Matches
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                                                                                                                       Query Match
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Best Local Similarity
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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                                                                              Local Similarity 100 les 8; Conservative
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                            VLRDDLLE 8
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                                                                           90.5%; Score 38; DB 4; L
100.0%; Pred. No. 1.7e+06;
Live 0; Mismatches 0;
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100.0%; Pred. No. 0.34;
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US-10-623-176-46

GENERAL INFORMATION:

Application US/10623176 No. US20040092446A1

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PRIOR FILING DATE: 2000-01-21
PRIOR PELING DATE: 1907-07-23
PRIOR FILING DATE: 1997-07-23
PRIOR PELING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR PELICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                    PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
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Best Local
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT PEPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/623,176 CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE: NAME/KEY: SITE
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ORGANISM: Artificial Sequence
                                                             TYPE: PRT
CRGANISM: Artificial Sequence
FEATURE: OTHEK INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                     LENGTH: 9
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8; Conserv
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100.0%; Pred. No. 1.7e+06;
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GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Engelhard, Victor H

TITLE OF INVENTION: The HA-1 Antigen

FILE REFERENCE: 2183-4285US

CURRENT APPLICATION NUMBER: US/10/791,217

CURRENT FILING DATE: 2004-03-02

PRIOR APPLICATION NUMBER: US/9/489,760

PRIOR APPLICATION NUMBER: US/99/489,760

PRIOR FILING DATE: 1998-07-23

PRIOR FILING DATE: 1998-07-23
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US-10-791-217-2
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; LOCATION: (1)...
US-10-623-176-2
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; OTHER INFORMATION: HA-1 peptide US-10-861-335-1
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                                                                                                                                                                                                 Sequence 1, Application US/10861335

Publication No. US20050031612A1

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa A.J.M.

TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunor

TITLE OF INVENTION: of tumors

FILE REFERENCE: 2183-6479US

CURRENT APPLICATION NUMBER: US/10/861,335

CURRENT FILING DATE: 2004-06-04

PRIOR APPLICATION NUMBER: PCT/NL02/00791

PRIOR APPLICATION NUMBER: EB 01204704.9

PRIOR APPLICATION NUMBER: ED 01204704.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 9
                                                                                                                    PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
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                                       ORGANISM: Artificial Sequence FEATURE:
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                                                                                   TYPE: PRT
                                                                                                     ENGTH:
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Local Similarity 88.9%;
es 8; Conservativo
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RESULT 15
US-10-623-176-43
(S-10-623-176-43)
; Sequence 43, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
; OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE: AME/KEY: SITE ; LOCATION: (1)..(10) US-10-623-176-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; SOPTWARE: PatentIn version 3.1
; SEQ ID NO 20
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments
US-11-007-740-20
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US-11-007-740-20
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                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/489,760
PRIOR PILING DATE: 2000-01-21
PRIOR PPLICATION NUMBER: EP 97202303.0
PRIOR PILING DATE: 1997-07-23
PRIOR PLICATION NUMBER: PCT/NL98/00424
PRIOR PILING DATE: 1998-07-23
PRIOR PPLICATION NUMBER: JP 2000-504165
PRIOR PILING DATE: 2000-12-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 1094-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 77
SOFTWARE: PatentIn Ver.
SEQ ID NO 43
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Best Local Similarity
Matches 8; Conserv
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATS: 2003-07-18
                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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Search completed: April 6, 2006, 17:28:45 Job time: 96.6667 secs
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                                                                               Matches
                                                                                        Best Local
                                                                                                 Query Match
                                      1 VLRDDLLEA 9
                                                                             Similarity
8; Conservat
                                                                             Conservative
                                                                                       88.1%;
                                                                             Score 37; DB 'Pred. No. 2.2; O; Mismatches
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB
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Perfect score:
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1: /SIDS5/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

2: /SIDS5/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /SIDS5/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

4: /SIDS5/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDS5/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

6: /SIDS5/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

7: /SIDS5/ptodata/2/pubpaa/US1_NEW_PUB.pep:*

8: /SIDS5/ptodata/2/pubpaa/US1_NEW_PUB.pep:*
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length: 15
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US-11-225-686-211
US-11-202-009-211
US-11-202-009-211
US-11-202-009-211
US-11-202-009-216
US-10-989-767A-219
US-10-989-767A-256
US-10-989-767A-268
US-10-989-767A-278
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169,
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45.2	45.2	47.6	47.6		47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6	47.6
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US-11-225-686-3877	US-11-225-686-2954	US-11-172-740-1404	US-11-126-841A-8	US-11-106-932-84	US-10-522-297-13	US-10-522-297-12	US-10-718-264-162	US-10-718-264-162	US-10-353-783-71	US-11-127-677-130	US-11-004-399-1909	US-11-254-419-67	US-11-254-419-60	US-10-927-634-63	US-10-927-435-63	US-11-021-305-66	US-11-021-305-65	US-11-021-305-62	US-11-096-706-164
Sequence 3877, Ap	Sequence 2954, Ap	Sequence 1404, Ap	Sequence 8, Appli	Sequence 84, Appl	Sequence 13, Appl	Sequence 12, Appl	Sequence 162, App	Sequence 162, App	Sequence 71, Appl	Sequence 130, App	Sequence 1909, Ap	Sequence 67, Appl	Sequence 60, Appl	Sequence 63, Appl	Sequence 63, Appl	Sequence 66, Appl	•	Sequence 62, Appl	Sequence 164, App

ALIGNMENTS

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RESULT 2
US-11-010-748A-11
; Sequence 11, Application US/11010748A
; Publication No. US20050244421A1
; GENERAL INFORMATION:
; APPLICANT: Merck Patent GmbH
; APPLICANT: STRITYMATTER, Wolfgang
; APPLICANT: MOLL, Heidrun
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US-11-010-748A-12
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Best Local S
Matches 9
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Publication No. US20050244421A1
GENERAL INFORMATION:
APPLICANT: Merck Patent GmbH
APPLICANT: STRITTMATTER, Wolfgang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.1 SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: MOLL, Heidrun
APPLICANT: SCHARM, Burkhard
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE
FILE REFERENCE: MER 136
CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
PRIOR APPLICATION NUMBER: PCT/EP03/06251
PRIOR PILING DATE: 2003-06-13
PRIOR PILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER: OF SEQ ID NOS: 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)..(9)
                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by TOTHER INFORMATION: ell epitopes
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SCHARM, Burkhard
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US-11-06-706-52

; Sequence 52, Application US/11096706

; Publicatton No. US20050245476A1
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Best Local S
Matches 4
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PA
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PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
SOPTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ZHOU, Hongxing
APPLICANT: ZHOU, Chen
APPLICANT: COSMAN, David J.
APPLICANT: CARTER, Paul
APPLICANT: CARTER, Paul
APPLICANT: MARTIN, Francis H.
APPLICANT: MARTIN, METHODS OF IDENTIFYING FUNCTIONAL ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: MER-136
CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/605,902
PRIOR FILING DATE: 2004-08-31
PRIOR APPLICATION NUMBER: 60/531,714
PRIOR FILING DATE: 2003-12-22
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: A-890A
CURRENT APPLICATION NUMBER: US/11/019,027
CURRENT FILING DATE: 2004-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: MISC_FEATURE
LOCATION: (1)..(9)
OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T
OTHER INFORMATION: ell epitopes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
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100.0%; Pred. No. 1.4e+05;
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Pred. No. 1.4e
0; Mismatches
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RESULT 6
US-11-25-686-211
; Sequence 211, Application US/11:
; Publication No. US20060019349A1
; GENERAL INFORMATION:
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US-11-225-686-208
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SEQ ID NO 208
LENGTH: 7
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APPLICANT: LIU, Qia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS FILE REFERENCE: 8325-0011.20 / S11-US2 CURRENT APPLICATION NUMBER: US/11/225,686 CURRENT FILING DATE: 2005-09-12 PRIOR FILING DATE: 2005-09-12 PRIOR FILING DATE: 2005-09-19 PRIOR FILING DATE: 2001-01-20 NUMBER OF SEQ ID NOS: 4085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sangamo Biosciences, Inc.
APPLICANT: Collingwood, Trevor
TITLE OP INVENTION: Treatment of Neuropathic Pain with Zinc Finger Proteins
FILE REFERENCE: 019496-008220US
CURRENT APPLICATION NUMBER: US/11/096,706
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 60/560,535
PRIOR FILING DATE: 2004-04-08
PRIOR PILING DATE: 2004-06-02
NUMBER: OF SEQ ID NOS: 227
RUGBER OF SEQ ID NOS: 227
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/225,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial FEATURE: OTHER INFORMATION: Chemically synthesized peptide (zinc finger protein recognition OTHER INFORMATION: region)
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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o. US20060019349A1
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Pred. No.
                                                                                                                                                                                                                                                                                                             Mismatches
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                                                             GNN NUCLEOTIDE
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RESULT 8
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US-11-225-686-216
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US-11-225-686-216
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                                                        GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: LIU Qiang

TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE

TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS

FILE REFERENCE: 8325-0011.20 / S11-US2

CURRENT PELICATION NUMBER: US/11/202,009

CURRENT FILING DATE: 2005-08-11

PRIOR APPLICATION NUMBER: US/999994

PRIOR FILING DATE: 2001-11-20

NUMBER OF SEQ ID NOS: 4085

SOFTWARE: PATENTIN Ver. 2.0

SEQ ID NO 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT FILING DATE: 2005-09-12
PRIOR APPLICATION NUMBER: US/91/989,994
PRIOR APPLICATION NUMBER: US/99/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOPTWARE: Patentin Ver. 2.0
LENGTH. 7
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Best Local S
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Publication No. US20060019349A1
                                                                                                                                                                                                                                                                                               Sequence 208, Application US/11202009 Publication No. US20060024726A1
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
PEATURE:
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4; Conserv
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Pred. No. 1.4e+05;
1; Mismatches 1
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Pred. No. 1.4e+05;
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Sequence 216, Application US/11202009

Publication No. US20060024726A1

GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: FRIPLETS BY ZINC FINGERS
FILE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/9/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOPTWARE: PATCENTIN Ver. 2.0
SEQ ID NO 216
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US-11-202-009-216
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; OTHER INFORMATION: Description of Artificial Sequence: example
US-11-202-009-208
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US-11-202-009-211
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REPERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PALENTIN Ver. 2.0
SEQ ID NO 211
LENGTH: 7
                                                   Query Match
Best Local
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Publication No. US20060024726A1
GENERAL INFORMATION:
                                  Matches
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Best Local Similarity
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                                                                                                                                            LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                               FEATURE: OTHER INFORMATION: Description of Artificial Sequence: example
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                                Local Similarity
nes 4; Conserv
3 RDDLLE 8
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                                  Conservative
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                                Score 21; DB 7;
Pred. No. 1.4e+05;
1; Mismatches 1
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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
ITILE OF INVENTION: UNCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158PID7
ITILE OF INVENTION: UNEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
ITILE OF INVENTION: OTHER CANCERS
ITLE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-04-10
NUMBER: 05/227,098
PRIOR FILING DATE: 2001-04-10
SOFTWARE: PAtentin Ver. 2.1
SEQ ID NO 219
LENGTH: 9
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US-10-989-767A-565
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APPLICANT: HUBERT, RENE
APPLICANT: HAITANO, ARTHUR
APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
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Matches
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Best Local Similarity
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APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: WICEBIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: OTHER CANCERS
FILE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR FILING DATE: 2000-08-22
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RAITANO, ARTHUR
AFAR, DANIEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LEVIN, ELANA
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Pred. No. 1.4e+05;
4; Mismatches 1
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APPLICANT: LEVIN, ELANA
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51582005004
CURRENT APPLICATION NUMBER: U$/10/989,767A
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 00/277,292
PRIOR APPLICATION NUMBER: 00/295,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR PILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-08-22
PRIOR PILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-989-767A-159
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RESULT 14
US-10-989-767A-268
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PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 565
LENGTH: 9
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Local Similarity 37.5%;
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Pred. No. 1.4e+05;
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Pred. No. 1.7e+02;
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GENERAL INFORMATION:
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE

Sequence 268, Application US/10989767A Publication No. US20060018917A1

APPLICANT: APPLICANT: APPLICANT:

LEVIN, ELANA CHALLITA-EID, PIA RAITANO, ARTHUR

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APPLICANT: RAITANO, ARTHUR
APPLICANT: RAITANO, ARTHUR
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLINA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: OFFICE AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OFFICE ANCERS
FILE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 00/277,292
PRIOR FILING DATE: 2001-08-12
PRIOR PILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 272
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
OTHER INFORMATION: Chemically synthesized peptide motif
US-10-989-767A-272
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FITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7

FITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND

FILE REFERENCE: 511582005004

CURRENT APPLICATION NUMBER: US/10/989,767A

CURRENT PRILING DATE: 2004-11-15

PRIOR APPLICATION NUMBER: US/277,292

PRIOR APPLICATION NUMBER: 09/935,430

PRIOR FILING DATE: 2002-10-21

PRIOR APPLICATION NUMBER: 09/935,430

PRIOR APPLICATION NUMBER: 60/227,098

PRIOR FILING DATE: 2001-08-22

PRIOR APPLICATION NUMBER: 60/282,739

PRIOR FILING DATE: 2001-04-10

SOPTWARE: Patentin Ver. 2.1

SEQ ID NO 268
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OTHER INFORMATION: Chemically synthesized peptide motif US-10-989-767A-268
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US-10-989-767A-272
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RAITANO, ARTHUR
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Pred. No. 1.7e+02;
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AAW99195 standard; peptide; 9 AA.

ALIGNMENTS

AAW99195;

20-MAY-1999 (first entry)

Minor histocompatibility antigen HA-1 T-cell epitope #1

Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.

Homo sapiens. Synthetic.

Misc-difference /label= His, Arg Location/Qualifiers

WO9905174-A1

04-FEB-1999.

23-JUL-1998; 98WO-NL000425

23-JUL-1997; 97EP-00202303.

(UYLE-) RIJKSUNIV LEIDEN.

Goulmy EAJM, Hunt DF, Engelhard €

WPI; 1999-153312/13.

A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation. HA-1 useful graft d l to treat : immune n bone

Claim 1; Page 32; 47pp; English.

RESULT 1
AAW991.95
ID AAW9
XX AAW
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XX Mino
XX M The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompactibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the

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RESULT 3
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                                                                                                                                                                                                                                                                                                  A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
 AAW99197 standard; peptide; 9 AA.
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Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9
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                                                                                                                T-cell epitope; minor histocompatibility antigen transplant rejection; Graft-versus-Host Disease; neoplastic haematopoietic cell.
                                                                                                                                                                                                      T-cell epitope
                                                                                                                                                                                                                                                                                                                               AAW97572 standard;
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                                                                                                                                                                                                                                              20-MAY-1999
                                                                                                                                                                                                                                                                                       AAW97572;
                   Misc-difference
                                                                            Homo sapiens
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                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                             peptide;
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"His or Arg"
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Pred. No.
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2e+06;
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                                                                                                                                              autoimmune disease;
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RESULT 5
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Neoplastic haematopoietic cells presenting the peptides, in an HIA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-and T-cells
Typing minor histocompatibility antigen HA-1 -
                      WPI; 1999-142960/12
                                                                                          23-JUL-1997;
02-JUN-1998;
                                                                                                                            23-JUL-1998;
                                                                                                                                                   04-FEB-1999
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                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                    Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                   HA-1 H-allele sequence.
                                                                                                                                                                                                                                                                                            13-MAY-1999
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                                                                      (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunogenic peptide from minor histocompatibility antigen HA-1 - for inducing tolerance to transplants and prevent rejection or {\sf gx}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYLE-) RIJKSUNIV LEIDEN
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                                                                                                                                                                                                                                                                                                                                        standard;
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                                                                                          97EP-00202303.
98EP-00870125.
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Pred. No.
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2e+06;
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n or graft-
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RESULT 6
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Best Local
The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used to screen for the HA-1 alleles. The HA-1 peptides can be used to screen for the HA-1 alleles. The HA-1 peptides can be used to screen for the HA-1 alleles. The HA-1 peptides can be used to screen for the HA-1 alleles. The HA-1 peptides can be used to screen for the HA-1 alleles.
                                                                                                                                                                            Claim 13;
                                                                                                                                                                                                             of genetic
                                                                                                                                                                                                                           Typing minor histocompatibility antigen HA-1 - identifying HA-1 R and HA-1 H alleles of HA-1,
                                                                                                                                                                                                                                                                                  WPI; 1999-142960/12
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02-JUN-1998;
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                                                                                                                                                                       English.
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Pred. No. 2e+0
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                                                                                               The invention relates to a novel method for providing epitopes of allelic covariants of antigenic proteins from specific species based on single concleotide polymorphism (SNP), by defining target protein/peptide or its concleoting allelic peptide/protein variants, expression product or its selecting allelic peptide/protein variants, expression product or its concerning epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is conserved proteins by applying the method for several protein from the concerning epitopes by profile of one or more individuals from a concerning epitopes binding to MHC protein. A protein of the invention has conserved the symplying the method for several protein from the concerning the supplying the method for several protein from the concerning the supplying the method for several protein from the concerning the supplying the method for several protein from the concerning the supplying profile was related to disease, preferably cancer This is useful for diagnosing a disease in an individual by concerning the supplying thematopoietic stem cells from a donor to a conserved profile that the invention concerning the supplying thematopoietic stem cells from a donor to a recipient and treating cancer, preferably leukaemia, and for determining the progression, regression or onset of a treated disease. The present extra concerning the supplication of the invention.
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                                                                       Sequence
                  Local Similarity
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                    97.3%;
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   Score 36; DB
Pred. No. 2e+0
0; Mismatches
   0;
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                    DB 8;
2e+06;
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     1;
                                      Length 9;
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       Indels
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       0,
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       Gaps
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ADT73023 ID ADT

ADT73023 standard; peptide; 9 AA.

RESULT 9

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1 VLXDDLLEA

Conservative

0

Mismatches

Indels

0,

Gaps

0,

2e+06;

VLRDDLLEA

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RESULT 8
ADH40334
ADH40334
AC ADH4
XX 
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Query Match
Best Local Similarity
Thes 8; Conservi
                                                                                                                                                                                                                                   The invention relates to a novel method for providing epitopes of allelic CC variants of antigenic proteins from specific species based on single CC nucleotide polymorphism (SNP), by defining target protein/peptide or its CR subset, screening database of DNA encoding target protein, identifying, CC selecting allelic peptide/protein variants, expression product or its CR fragment encoded by DNA sequence having SNP, creating variant epitopes, CC selecting epitopes binding to MHC protein. A protein of the invention has CR cytostatic activity, and may have a use in a vaccine. The method is CR useful for generating a SNP profile of one or more individuals from a CR civilate where the SNP profile of one several protein from the CR individuals, where the SNP profile was related to disease, preferably CR cancer. This is useful for diagnosing a disease in an individual by CR generating the SNP-related polymorphic profile. A method of the invention CR is useful for transplanting haematopoietic stem cells from a donor to a ceipient and treating cancer, preferably leukaemia, and for determining CR the progression, regression or onset of a treated disease. The present CR sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human minor histocomptibility antigen HA-1 T cell eptiope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-JUN-2002; 2002EP-00013423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUN-2003; 2003WO-EP006251
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                                                                                                                                                                             Sequence 9 AA;
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                                                                     97.3%;
                                                                         Score 36;
Pred. No.
                                                                                                               DB 8;
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RESULT 10
ABY01546
ID ABY01
XX
AC ABY01
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DT 16-JU
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DE SARS
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                                                                                                                                                                                                                                                                                                                                                                 CC This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein I and CC sight structural proteins preferably G also M, M2-1, M2-2, N, P, F, and CC SH (also known as the 1A protein). Specifically, it refers to a CC composition comprising an above peptide mixed with a pharmaceutical cc excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC Tlymphocytes (CTLs) that respond to a major histocompatibility complex (CMHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (prophylactic or therapeutic) vaccine composition for a DNA-based CC immunisation, or for preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the CC polypeptide in an immune response provoking an activity. This CC peptide sequence is a human RSV high binding affinity peptide of the CC invention.
                                                                                                                                                                                                                                                                        Query Match
Best Local (
SARS coronavirus spike protein HLA A 0201 T-cell epitope, SEQ:8174.
                                   16-JUN-2005
                                                                    ABY01546;
                                                                                                    ABY01546 standard;
                                                                                                                                                                                                                                                                                                                        Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lasters I, Desmet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-2003; 2003EP-00447095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human respiratory syncytial virus; cytotoxic T lymphocyte; CTL; MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
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                                                                                                                                                                                                                                                     Local Similarity
es 4; Conserv
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                                 (first entry)
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Pred. No. 2e+0
3; Mismatches
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2e+06;
                                                                                                                                                                                                                                                                                       Length 9;
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Vaccine; nucleic acid vaccine; drug screening; diagnosis; SARS coronavirus infection; infection; respiratory disease; virucide. SARS coronavirus.

WO2004092360-A2.

28-OCT-2004.

09-APR-2004; 2004WO-US011710.

10-APR-2003; 2003US-046218P.
11-APR-2003; 2003US-0462418P.
11-APR-2003; 2003US-0462418P.
11-APR-2003; 2003US-046310P.
11-APR-2003; 2003US-046310P.
11-APR-2003; 2003US-046310P.
11-APR-2003; 2003US-046310P.
11-APR-2003; 2003US-046393P.
11-APR-2003; 2003US-046393P.
12-APR-2003; 2003US-046899P.
22-APR-2003; 2003US-046899P.
22-APR-2003; 2003US-046899P.
23-APR-2003; 2003US-046899P.
23-APR-2003; 2003US-046899P.
23-APR-2003; 2003US-0468912P.
21-APR-2003; 2003US-0468912P.
21-APR-2004; 2004US-0566757P.
(CHIR ) CHIRON CORP.
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Rappuoli R, Masignani V, Stadler K, Gregersen J, Chien D, H Polo J, Weiner A, Houghton M, Song HC, Seo MY, Donnelly JJ; Klenk HD, Valiante N;

PI; 2004-766863/75

Novel isolated polypeptide e.g. spike polypeptide, Env polypeptide, of severe acute respiratory syndrome virus (SARS), useful as vaccine for SARS.

Disclosure; SEQ ID NO 8174; 839pp; English.

The invention relates to isolated polypeptides of the severe acute CC respiratory syndrome (SARS) coronavirus. The polypeptides include spike CC (S or E2), env (E or sM), membrane (M or E1), hemagglutinin-esterase (HE CC or E3), and nucleocapsid (N) polypeptides, and the ORF1a and ORF1ab CC (replicase) polypeptides and their proteolytic fragments. The invention CC also relates to antibodies which recognise the polypeptides; nucleic codis encoding the SARS virus polypeptides; primers specific for SARS virus nucleic acids sequences; kits for amplifying SARS virus target nucleic acids; a double-stranded RNA molecule 10-30 nucleotides in length CC which is able to inactivate the SARS virus in a mammalian cell; an CC expression construct for recombinant expression of a SARS virus spike CC protein; a viral vector for in vivo delivery of a SARS virus polypeptide-cc encoding nucleic acid; and a mammalian cell line stably expressing a SARS virus antigen. The invention additionally provides a vaccine for the treatment or prevention of SARS comprising an inactivated SARS virus, a split SARS virus, a split SARS virus, a cc killed SARS virus, an attenuated SARS virus, a split SARS virus, a cc replicon particle comprising one or more SARS virus antigens; methods of comparatory virus antigens; method soff constitying a therapeutically active agent by measuring the effect of the agent on a SARS-related enzyme, and a method of treating a SARS patient comparation and inhibitors. The SARS virus polypeptides and colded caids can be used in the preparation and manufacture of vaccines

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RESULT 11
The invention relates to a cytotoxic Yersinia pestis T-cell epitope, considered from any of the 1000 sequences of 9 amino acids appearing as ADZ49933-ADZ50932. Also included are predicting peptides that are captions or can be used as diagnostic tools (comprising predicting which peptides bind to a HMC molecule (not defined) with high affinity using a neural network with at least one of the following features: some or all contents of the inputs to the neural networks are generated using a hidden Markov model; or some or all of the inputs are encoded by an amino acid substitution matrix, different from an identity matrix) and a vaccine or diagnostic tool using a limited number such as at least 1, 2, 3, 4, 5, 8, 16, 32, 64, 128, 256, 512 of the peptides of ADZ49933-ADZ50932. In predicting peptides, the prediction of the neural network is combined with prediction or measurement of one of the following: proteasomal cleavage sites; MHC binding; presence of sequence or related sequence(s) in patent databases; TAP binding; gene or protein expression level; function of the protein; localization of the protein; and similarity to self proteins. The epitope is useful as a vaccine or diagnostic tool, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 9
                                                                                                                                                                                                                                                                                                                                                                       in a
                                                                                                                                                                                                                                                                                                                                                                                            New cytotoxic Yersinia useful as a vaccine or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yersinia pestis infection; antibacterial; infection; epitope; T-cell; vaccine; biological warfare.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Y. pestis cytotoxic T-cell
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                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 993; 235pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nielsen
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Pred. No. 2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                          10 amino acids, immune response
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RESULT 12
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for inducing immune response in a subject (said immunity to yersinia pestis infection, the causative agent of plague). The present sequent a cytotoxic Yersinia pestis T-cell epitope of the invention.
                                                      sequences AAU71093-AAU71796 represent the 103P3E8-related protein and peptide fragments of the protein. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 polynucleotide, its related protein and peptide fragments and specific PCR primers are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P3E8-related protein, and a ribozyme capable of cleaving a polynucleotide having the 103P3E8 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P3E8. The sequences can be used in diagnostic methods to monitor the level of 103P3E8 gene products in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          103P3E8; prostate; bladder; kidney; colon; lung; breast; rectum; stomach; tumour; cancer; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human;
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                                                                                                                                                                                                                                                                                    Disclosure; Page 84; 128pp; English.
                                                                                                                                                                                                                                                                                                                                 Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, exposes cancer, comprises determining presence of aberrant 103P3E8 generate cancer,
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                             cancerous
                                            serum, blood,
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                                                                                                                                                                 Sequences AAU71093-AAU71796 represent the 103P3E8-related protein and peptide fragments of the protein. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 polymucleotide, its related protein and peptide fragments and specific PCR primers are therefore useful for diagnosing and treathing cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P3E8-related protein, and a ribozyme capable of cleaving a polymucleotide having the 103P3E8 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P3E8. The sequences can be used in diagnostic methods to monitor the level of 103P3E8 gene products in serum, blood, urine and tissue and to thereby detect the presence of
                                                                                                                           Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 90; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene
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                                                                                                                                                          cancerous
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                                                           62.2%; Score 23; DB
100.0%; Pred. No. 2e-
cive 0; Mismatches
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CC polypeptide in an immune response-provoking affinity peptide of the immunical activity. This composition is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                            Query Match
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ADT72767;
                                                                ADT72767 standard; peptide; 9 AA.
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Human RSV L high affinity binding peptide SeqID 663

13-JAN-2005

(first entry)

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                                                                                                                                          Query Match
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Matches 4
                                                                                                                                                                                                                 Sequence 9 AA;
                                                                                                                                                                                                                                                                  polypeptide in an immune response-provoking amount of specific CTL). Accordingly, these peptide compositions have virucidal activity. This peptide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 663; 143pp; English.
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: pir2:*
3: pir3:*
4: pir4:*
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46.390 Million cell updates/sec
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A59489

A61328

PH0942

PH0942

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PT06601

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A34026

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PT0722

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cerebellar degener
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calsequestrin, fas
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elastase - Pseudom
protein kinase C i
                     ferredoxin a2 - Ja
tetrameric protein
tetrameric protein
chlorophyll a/b-bi
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                                                                158K exoantigen
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S

hucolin, 75K chain - human (fragment) c;Species: Homo sapiens (man) C;Date: 15-Feb-1997 #sequence_revisio: C;Accession: 868004 R;Edgar, P.F. A;Status: preliminary A;Molecule type: protein A;Residues: 1-7 <EDG> PEBS Lett. 375, 159-161, 1995 A;Title: Hucollin, a new cortic A;Reference number: \$68004; M A;Accession: \$68004 A;Cross-references: UNIPARC:UPI000017C164 Query Match Best Local S Matches 3 4 DDL 6 Similarity 3; Conserv Conservative new corticosteroid-binding protein from S68004; MUID:96087107; PMID:7498469 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999 43.2%; Score 16; DB 2; I 100.0%; Pred. No. 2.8e+05; tive 0; Mismatches 0; 0, Length 7; human 0; plasma with structural Gaps 0

RESULT S68004

ALIGNMENTS

R;KawaBaki, S.; Ārai, H.; Igarashi, Y.; Kodama, T.

Gene 167, 87-91, 1995
A;Title: Sequencing and characterization of the downstream region of the genes encod:
y for biosynthesis of heme dl.
A;Reference number: JC4552; MUID:96144254; PMID:8566817
A;Accession: PC4131
A;Status: preliminary
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-8 <KAW>
A;Cross-references: UNIPROT:P95412; UNIPARC:UPI000017A96A; DDBJ:D50473; NID:g1217594
A;Note: this ORF is not annotated in GenBark entry PSENIRC, release 113.0 ঠ A;Note: this ORF is not annotated in Gene C;Superfamily: Pseudomonas stutzeri nirD hypothetical protein 8 [imported] - Pseudomonas aeruginosa (fragment) C;Species: Pseudomonas aeruginosa C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 31-Dec-2004 C;Accession: PC4131 밁 Best Loc Matches Query Match Local Similarity
3; Conserv 9 Tag Conservative 0; 43.2%; Score 16; Pred. No. Mismatches protein DB 2; Lo 0; Length 8; Indels region of the genes encoding <u>,,</u> Gaps 0,

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A;Description: Purification A;Reference number: A59489 A;Accession: A59489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein kinase C inhibitor - rat (fragment) C;Species: Rattus norvegicus C;Date: 25-Aug-2003 #sequence_revision 25-AC;Accession: A59489
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FEBS Lett. 299, 291-293, 1992
A;Title: Identification of cleavage sites involved in proteolytic processing
A;Reference number: S20446; MUID:92183956; PMID:1544509
A;Accession: S20446
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      elastase - Pseudomonas aeruginosa
C;Species: Pseudomonas aeruginosa
C;Date: 19-Mar-1997 #sequence_revision 24-Jul-1997 #text_change 24-Jul-1997
                                                                                    A;Cross-references: UNIPROT:Q7M390; UNIPARC:UPI000017C43F
C;Keywords: hydrolase; protein digestion; serine proteinase; zymogen
F;1-8/Domain: activation peptide #status experimental <APT>
                                                                                                                                                                                                                                                                                              trypsin (EC 3.4.21.4) precursor - minke whale (fragment)
C;Species: Balaenoptera acutorostrata (minke whale, lesser rorqual)
C;Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004
C;Accession: A61328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: protein
A; Residues: 1-7 <NEG>
A; Experimental source: strain Wistar, li
A; Note: p-Hydroxyacetophenone-Sepharose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        submitted to the Protein Sequence Database, June 2003 A; Description: Purification of PKCI from rat liver.
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                                                                                                                                                                                  A;Reference number: A61328; MUID:75146765; PMID:1125273 A;Accession: A61328
                                                                                                                                                                                                                                                         R;Bricteux-Gregoire, S.; Schyns, R.; Florkin, M.;
Biochim. Biophys. Acta 386, 244-255, 1975
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                               35.1%;
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Pred. No. 2.8e
2; Mismatches
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Pred. No. 2.8e+05;
                               Score 13; DB 2;
Pred. No. 2.8e+05;
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2.8e+05;
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                                                                                      R;Shiuan, D.; Campbell, A. Gene 67, 203-211, 1988
A;Title: Transcriptional regulation and gene arrangement A;Reference number: I40697; MUID:89006280; PMID:2971595
                                                                                                                                                              biotin A - Citrobacter freundii (fragment)
C;Species: Citrobacter freundii
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C;Accession: 140697
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-9 < GOL>
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J. Exp. Med. 174, 1467-1476,
A;Title: Analysis of T cell J
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               A;Status: preliminary; translated from GB/EMBL/DDBJA;Molecule type: DNAA;Residues: 1-4 <RES>
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of.

Escherichia coli,

Citrobacte

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macrophage cytotoxicity-inducing factor, 29K - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
C;Accession: A60427
R;Jones, C.M.; Prince, C.A.; Williams, J.S.
Exp. Hematol. 19, 704-709, 1991
A;Title: Purification and amino acid analysis of a human macrophage cytotoxicity-inducin A;Heference number: A60427; MUID:91372335; PMID:1909970
A;Accession: A60427
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A;Acession: PH0942
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C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH0942
C;Accession: PH0942
                                                                                                   A;Molecule type: protein
A;Residues: 1-9 <JON>
A;Residues: 1-9 <JON>
A;Rosidues: 1-9 <INIPROT:Q7M4R5; UNIPARC:UPI0000142BE0
A;Cross-references: UNIPROT:Q7M4R5; UNIPARC:UPI0000142BE0
C;Note: the sequence from the text on page 706 is inconsistent with that from page 708
C;Keywords: cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell receptor beta chain V-D-J region (clone 13) - rat
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A; Cross-references:

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J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell receptor beta chain V-D-J region (120-2K) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 17-Jul-1992 #sequence revision 17-Jul-1992 #text_change 09-Jul-2004 C;Accession: PT0601; PT0617; PT0694
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PT0679
A;Cross-references: UNIPARC:UPI000017C7E6
A;Experimental source: day 18 fetal thymus,
C;Keywords: T-cell receptor
                                                                     A; Molecule type: DNA
A; Residues: 1-5 < FE2>
                                                                                                                                                                                               A;Status: translation not shown A;Molecule type: mRNA A;Residues: 1-5 <PE3>
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0679
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0679; PT0708
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A;Experimental source: newborn thymus, strain BALB/c,
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A;Experimental source: newborn thymus, strain BALB/c, clone 120-2K
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A; Residues: 1-5 < FEE>
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A;Experimental source: newborn thymus, strain BALB/c, C;Keywords: T-cell receptor
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A; Residues: 1-5 < FE2>
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A;Experimental source: day 18 fetal thymus,
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100.0%; Pred. No. 2.8e+05;
tive 0; Mismatches 0;
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acetylcholinesterase (EC 3.1.1.7) 5.6S form - Pacific electric ray (fragment) c;Species: Torpedo californica (Pacific electric ray)
C;Date: 16-Mar-1990 #sequence_revision 16-Mar-1990 #text_change 08-Nov-1996
C;Accession: A34026
R;Gibney, G.; MacPhee-Quigley, K.; Thompson, B.; Vedvick, T.; Low, M.G.; Taylor, S.S.; '1
J. Biol. Chem. 263, 1140-1145, 1988
A;Title: Divergence in primary structure between the molecular forms of acetylcholinests A;Reference number: A34026; MUID:88087239; PMID:3335534
A;Accession: A34026
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J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T ce
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C;Date: 28-Sep-1990 #sequence_revision 28-Sep-1990 #text_change 24-Jun-1993
C;Accession: B35640
R;Chen, Y.T.; Rettig, W.J.; Yenamandra, A.K.; Kozak, C.A.; Chaganti, R.S.K.; Posner, Proc. Natl. Acad. Sci. U.S.A. 87, 3077-3081, 1990
A;Title: Cerebellar degeneration-related antigen: a highly conserved neuroectodermal A;Reference number: A35640; MUID:90222173; pMID:2326268
A;Accession: B35640
A;Accession: B35640
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A; Residues: 1-6 < FEE>
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A; Residues: 1-6 < CHE>
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calsequestrin, fast skeletal muscle - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 23-Aug-1991 #sequence_revision 23-Aug-1991 #text_change 23-Feb-1997
C;Accession: B39040
R;Cala, S.E.; Jones, L.R.
J. Biol. Chem. 266, 391-398, 1991
A;Hiol. Phosphorylation of cardiac and skeletal muscle calsequestrin isoforms by casein A;Reference number: A39040; MUID:91093153; PMID:1985907
A;Accession: B39040
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(;Species: Mus musculus (house mouse)
(;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
(;Accession: PT0628
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0628
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A;Residues: 1-7 <GIB>
A;Cross-references: UNIPARC:UPI000017BF32
C;Keywords: alternative splicing; carboxylic ester hydrolase
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A;Molecule type: mRNA
A;Residues: 1-7 <FEB>
A;Cross-references: UNIPARC:UPI000017C7CD
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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A;Molecule type: protein
A;Residues: 1-7 <CAL>
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Search completed: April 6, 2006, 16:26:10 Job time: 20.6667 secs
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Best Local Similarity '5...
"-rches 3; Conservative
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Best Local Similarity
Matches 2; Conserv
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100.0%; Pred. No. 2.8e+05;
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           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 0.5
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37
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Q70SM2_HUMAN
Q8LFF15_MAIZE
P82568_STRPY
Q9XJN0_9VIRU
Q9FSZZ_CICAR
Q9GYZAB_MOUSE
Q85710_9RETR
Q8UTD7_9HIV1
C99182_SYMEGE
Q6UTR2_CRYNV
Q9HDS4_ASPFL
Q7M390_BALAC
Q7M390_BALAC
Q7M390_BALAC
Q7M390_BACAC
Q7M390_BACAC
Q8H9K1_9CAUD
Q8SBJO_BPR69
Q6JCK8_SOYEN
Q6JCK8_SOYEN
Q72279_RHOGAN
Q727279_RHOGAN
Q9BQT4_HUMAN
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RESULT 2 QBLPT5_MAIZE QBLPT5_MAIZE PRELIMINARY; PRT; 9 AA. AC QBLPT5; DT 01-0CT-2002 (TrEMBLrel. 22, Created) DT 01-0CT-2002 (TrEMBLrel. 22, Last sequence update) DT 01-0CT-2002 (TrEMBLrel. 22, Last amoutation update) DE Beta-expansin-like protein (Fragment). CE ukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheop CE spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; CE PACCAD clade; Panicoideae; Andropogoneae; Zea. OX NCBI_TaxID=4577; RN 11 RP NUCLEOTIDE SEQUENCE. RA Ching A.S., Caldwell K.S., Jung M., Dolan M., Smith O.S., Tin RA Morgante M., Rafalski J.A.; RI Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AY094310; AAM21836.1; -; Genomic_DNA.	RESULT 1 Q70SM2_HUMAN ID Q70SM2_HUMAN PRELIMINARY; PRT; 9 AA. AC Q70SM2; DT Q5_JUL_2004 (TrEMBLrel. 27, Created) DT O5_JUL_2004 (TrEMBLrel. 27, Last sequence update) DT O5_JUL_2004 (TrEMBLrel. 27, Last annotation update) DT O5_JUL_2004 (TrEMBLrel. 27, Last annotation update) DF UPPermethylated in cancer 1 (Fragment). GN NOTER (Human). OC EMMANYOTA; Metazoa; Chordata; Craniata; Vertebrata; Euteleost Mammalla; Buarchontoglires; Primates; Catarrhini; H OC Homo. OX NOCEL TAXID=9606; RN NUCLEOTIDE SEQUENCE. RT TISSUB_Mammary gland; Pinte S., Guerardel C., Deltour S., Godwin A.K., Leprince D.; RA Pinte S., Guerardel C., Deltour S., Godwin A.K., Leprince D.; RA Pinte S., Guerardel C., Deltour S., Godwin A.K., Leprince D.; RA Pinte S., Guerardel C., Deltour S., Godwin A.K., Leprince D.; RT murine and rat tumor suppressor genes HIC1."; DC COMPANA AND SEQUENCE. RT MURING SEQUENCE SON SECUENCE SO	32 13 35.1 9 2 Q56SU1_9DIPS Q568u1 33 13 35.1 9 2 Q56SU3_9DIPS Q568u2 34 13 35.1 9 2 Q56SU3_9DIPS Q568u3 35 13 35.1 9 2 Q56SU3_9DIPS Q568u4 36 13 35.1 9 2 Q56SU4_9DIPS Q568u4 37 13 35.1 9 2 Q56SU5_9DIPS Q568u5 38 13 35.1 9 2 Q56SU5_9DIPS Q568u5 39 13 35.1 9 2 Q56SU5_9DIPS Q568u7 39 13 35.1 9 2 Q56SU7_9DIPS Q568u8 40 13 35.1 9 2 Q56SU9_9DIPS Q568u8 41 13 35.1 9 2 Q56SU9_VIBLN 42 13 35.1 9 2 Q56SV1_9DIPS Q568u8 43 13 35.1 9 2 Q56SV1_9DIPS Q568v1 44 13 35.1 9 2 Q56SV1_9DIPS Q568v1 45 13 35.1 9 2 Q56SV1_9DIPS Q568v1 46 13 35.1 9 2 Q56SV1_9DIPS Q568v1 47 13 35.1 9 2 Q56SV1_9DIPS Q568v2 48 13 35.1 9 2 Q56SV1_9DIPS Q568v2 49 13 35.1 9 2 Q56SV1_9DIPS Q568v2 40 13 35.1 9 2 Q56SV1_9DIPS Q568v2 41 13 35.1 9 2 Q56SV1_9DIPS Q568v2 42 13 35.1 9 2 Q56SV1_9DIPS Q568v2 43 13 35.1 9 2 Q56SV1_9DIPS Q568v2 44 13 35.1 9 2 Q56SV1_9DIPS Q568v2 45 13 35.1 9 2 Q56SV1_9DIPS Q568v3
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01-NOV-1999 (TrEMBLrel.
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Submitted (MAY-2000)
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01-OCT-2000 (TrEMBLrel. 15, Last annotat:
01-OCT-2003 (TrEMBLrel. 25, Last annotat:
Unknown protein from 2D-page (Fragment).
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                                                                                                                                                          SEQUENCE
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VanBogelen R.A.;
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                                                                                                                                                                                                                                                                                         Mindich L., Qiao X., Qiao
Hoogstraten D.;
                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
MEDLINE=99350412; PubMed=10419946;
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                                                                                                                                                                                                                                                                    "Isolation of additional bacteriophages
                                                                                                                                                                                                                                          louble-stranded RNA.";
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Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases
EMBL; AF192526; AAF04843.1; -; Genomic_DNA.
MGI; MGI:2136650; Clec2i.
MGI; MGI:2136650; Dcl1.
GO; GO:0009897; C:external side of plasma membrane; IDA
GO; GO:0005887; C:integral to plasma membrane; TAS:
GO; GO:0046703; F:natural killer cell lectin-like recep
GO; GO:0005515; F:protein binding; IPI.
GO; GO:0016066; P:cellular defense response (sensu Vert
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Submitted (SEP-2000) to the
EMBL; AJ299069; CAC10216.1;
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid
eurosids I; Fabales; Fabaceae; Papilionoideae; Cicereae; Cicer.
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01-OCT-2003 (TrEMBLrel. 25,
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Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognathi,
Muroidea, Muridae, Murinae, Mus.
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Name=Clec2i; Synonyms=Dcl1;
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DOI=10.1128/JVI.76.11.5435-5451.2002;
Novitsky V., Smith U.R., Gilbert P., McLane M.F., Chigwed Williamson C., Ndung'u T., Klein I., Chang S.-Y., Peter T Foley B.T., Gaolekwe S., Rybak N., Gaseitsiwe S., Vannber Marlink R., Lee T.-H., Essex M.;
"Human immunodeficiency virus type 1 subtype C molecular consensus sequence for an AIDS vaccine design?";
J. Virol. 76:5435-5451(2002).
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01-NOV-1996
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Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF443091; AAL34712.1; -; Genomic_DNA.
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MEDLINE-84115080; PubMed-6319754;
MEDLINE-84115080; PubMed-6319754;
Lerner T.L., Hanafusa H.;
"DNA sequence of the Bryan high-titer strain of Rous sarcoma virus:
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GN Name=COI;
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OG Mitochondrion.
OC Eukaryota; Met
OC Actinopterygii
OC Actinopterygii
OC Actinopterygii
OC Actinopterygii
OC Acyprinodontifo
OX NCBI TAXID=135
RP NUCLEOTIDE SEQ
MEDLINE=200729
RA Murphy W J., T
RT "Phylogeny of
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RT Sequences.";
RL Mol. Phylogene
DR EMBL; AF002591
DR GO; GO:0005739
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                                                                                                                                                                                                                                                                       Cryptococcus neoformans var. grubii H99. Eukaryota; Fungi; Basidiomycota; Hymenom; Tremellomycetidae; Tremellales; Tremellales; MCBI_TaxID=235443;
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NCBI_TaxID=135316;
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                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WEDLINE=20072928; PubMed=10603257; DOI=10.1006/mpev.1999.0656; Murphy W.J., Thomerson J.E., Collier G.E.; Phylogeny of the Neotropical killifish family Rivulidae (Cyprinodontiformes, Aplocheiloidei) inferred from mitochondri
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RESULT 12

Q7M390_BALAC

Q7M390_BALAC

Q7M390_BALAC

Q7M390;
DT 01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last sequence)
DT 01-MAR-2004 (TrEMBLrel. 26, Last sequence of trypsinogen from the rorqual, Balaenoptera acutorostrata (Cetacea). Simultan PROPERIOR (Trypsinogen, Chymotrypsinogen and ribonuclease from PROPERIOR (Trypsinogen, Chymotrypsinogen and ribonuclease from the rorqual, Balaenoptera acutorostrata (Cetacea). Simultan of trypsinogen, chymotrypsinogen and ribonuclease from PROPERIOR (Propsinogen, Chymotrypsinogen and ribonuclease from PROPERIOR (Propsinogen, Chymotrypsinogen and ribonuclease from PROPERIOR (Propsinogen, Chymotrypsinogen and Propsinogen, Chymotrypsinogen and Propsinogen Scales (Propsinogen, Chymotrypsinogen Scales (Propsinogen, Chymotrypsinogen Scales (Propsinogen, Chymotrypsinogen Scales (Propsinogen, Chymotrypsinogen Sc
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01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
TrpC polyprotein (Fragment)
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Q9HDS4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "N-terminal amino acid sequence of trypsinogen from the less rorqual, Balaenoptera acutorostrata (Cetacea). Simultaneous of trypsinogen, chymotrypsinogen and ribonuclease from pancz Biochim. Biophys. Acta 386:244-255(1975).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Cetacea;
Mysticeti; Balaenopteridae; Balaenoptera.
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trypsinogen, chymotrypsinogen and ribonuclease from pancreas.";
ochim. Biophys. Acta 386:244-255(1975).
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RESULT 13
QRH9K1 9CAUD PRELIMINARY;
ID QRH9K1;
AC QRH9K1;
AC QRH9K1;
AC QRH9K1;
AC QRH9K1;
DT 01-MAR-2003 (TrEMBLrel. 23,
DT 01-MAR-2003 (TrEMBLrel. 23,
DT 01-MAR-2003 (TrEMBLrel. 23,
DT 01-MAR-2003 (TrEMBLrel. 23,
DR GP30.2 protein (Fragment).
GN Name-30.2;
OS Bacteriophage LZ10.
OC Viruses; daDNA viruses, no
OC T4-like viruses,
OX NCBI TaxID-192973;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA KOlesinskiene G., Nivinskas
RL Submitted (APR-2002) to the
DR EMBL; AJ458400; CAD30256.1;
FT NON TER
SQ SEQUENCE 8 AA; 918 MW;
                                                                                    QRSBJO BPR69
ID QRSBJO B
AC QRSBJO B
AC QRSBJO C
DT 01-JUN-2
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CR NAME=30.
OS BACTETIC
OC VITUSES
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OX NCBI TAN
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                                                                                                    Nivinskas R.;
Submitted (MAR-2002) to the
Submitted (MAR-2002) to the
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                                                                                                                                                                          Viruses; dsDNA viruses,
T4-like viruses.
NCBI_TaxID=12353;
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Submitted (APR-2002) to the
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-; Genomic_DNA.
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No. 2
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RESULT 15
Q6JC68 SOYBN PRELIMINARY;
ID Q6JC68_SOYBN PRELIMINARY;

PRT;

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AC Q6JC68;
DT 05-UUL-2004 (TrEMBLrel. 27, Last sequence update)
DT 05-UUL-2004 (TrEMBLrel. 27, Last sequence update)
DT 05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
DE Isoflavone synthase 1 (Fragment).
OS Glycine max (Soybean).
OC Enkaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
OC Spermatophyta; Magnoliophyta; endicotyledone; core eudicots; rosids;
OC eurosids I; Fabales; Pabaceae; Papilionoideae; Phaseoleae; Glycine.
OX NCBI_TAXID=3847;
RN NUCLEOTIDE SEQUENCE.
RN PUMMed=1535384; DOI=10.1023/B:PLAN.0000040814.28507.35;
RN PUMMed=1535384; DOI=10.1023/B:PLAN.00000040814.28507.35;
RN PUMMed=1535384; DOI=10.1023/B:PLAN.000000040814.28507.35;
RN PUMMed=1535384; DOI=10.1023/B:PLAN.000000040814.28507.35;
RN PUMMed=
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Result
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Maximum DB
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Maximum Match
Listing first
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Perfect score:
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seq length: 9
              protein search, using sw model
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
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              572060 seqs, 82675679 residues
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Copyright (c) 1993 - 2006 Biocceleration
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US-08-556-419-14
US-10-394-980-464
US-08-459-688-20
US-08-399-411-72
US-08-516-859A-75
US-08-516-859A-75
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Sequence 18, Appl Sequence 29, Appl Sequence 29, Appl Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 6, Appli Sequence 14, Appl Sequence 14, Appl Sequence 421, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 75, Appl Sequence 76, Appl Sequence 77, Appl Sequence 78, Appl Sequence 78
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US-09-269-250E-20 US-09-269-250E-20 ; Sequence 20, Application US/09269250E ; Patent No. 6830883 ; GENERAL INFORMATION: APPLICANT: Goulmy, Elsa ; TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 ; FILE REPERSENCE: 58994 ; CURRENT APPLICATION NUMBER: US/09/269,250E ; CURRENT FILING DATE: 1999-05-21 ; NUMBER OF SEQ ID NOS: 38 ; SOFTWARE: Patentin version 3.1 ; SEQ ID NO 20 ; LENGTH: 9 ; TYPE: PRT ; ORGANISM: Artificial Sequence ; OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-09-269-250E-20	Query Match 97.3%; Score 36; DB 2; Length 9; Best Local Similarity 88.9%; Pred. No. 4.6e+05; Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0; Qy 1 VLXDDLLEA 9	RESULT 1 US-09-269-250E-18 ; Sequence 18, Application US/09269250E ; Sequence 18, Application US/09269250E ; Patent No. 6830883 ; Patent Polication Number: US/09/269,250E ; CURRENT FILING DATE: 1999-05-21 ; NUMBER OF SEQ ID NOS: 38 ; SOFTWARE: Patent Nors: 38 ; SOFTWARE: Patent Nors: 38 ; Pature: 199-05-21 ; ORGANISM: Artificial Sequence ; Pature: 199-269-250E-18	28 21 56.8 9 2 US-09-217-609A-11 Sequence 11, Appl 29 21 56.8 9 2 US-08-873-235B-11 Sequence 11, Appl 30 21 56.8 9 2 US-09-586-472-72 Sequence 72, Appl 31 21 56.8 9 2 US-09-586-472-78 Sequence 78, Appl 32 21 56.8 9 2 US-09-528-706-72 Sequence 78, Appl 34 21 56.8 9 2 US-09-528-706-72 Sequence 78, Appl 35 20 54.1 6 2 US-09-025-819-11 Sequence 11, Appl 36 20 54.1 6 2 US-09-025-819-11 Sequence 11, Appl 37 20 54.1 6 2 US-09-808-126-11 Sequence 11, Appl 39 20 54.1 8 2 US-08-803-951-11 Sequence 11, Appl 39 20 54.1 8 2 US-08-747-599A-79 Sequence 11, Appl 43 19 51.4 6 2 US-08-747-599A-50 Sequence 50, Appl 43 19 51.4 6 2 US-08-974-549A-73 Sequence 373, Appl 44 19 51.4 6 2 US-09-402-181B-373 Sequence 373, Appl 51.4 6 2 US-09-721-456-373 Sequence 373, Appl 51.4 6 2 US-09-721-456-373 Sequence 373, Appl 51.4 7 2 US-09-187-859-1186
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Query Match

97.3%;

Score 36; DB 2;

Length 9;

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Best Local Matches

Similarity 88.9%; 8; Conservative

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Pred. No. 4.6e+05; 0; Mismatches 1;

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Gaps

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US-09-269-250E-29

Sequence 29, Application Patent No. 6830883 GENERAL INFORMATION:

US/09269250E

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FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILLING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILLING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
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APPLICANT: GOLINY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN Version 3.1
SEQ ID NO 29
LENGTH: 9
TYPE: PRT
ORGANISM: Human
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (3)..(3)
OTHER INFORMATION: Xaa represents a histidine (H) or an arginine (R) residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
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                                                                           Query Match
Best Local &
                                                          Matches
                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Histocompatibility antigen FEATURE:
                                                                                                                                                      NAME/KEY: PEPTIDE LOCATION: (3)
LOCATION: (3)
OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE OTHER INFORMATION: RESIDUE
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1 VLXDDLLEA 9
                                                      h 97.3%; Score 36; DB 2; L
Similarity 100.0%; Pred. No. 4.6e+05;
9; Conservative 0; Mismatches 0;
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; Sequence 38, Application ; Patent No. 6830883 ; GENERAL INFORMATION:

US/09269250E

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APPLICANT: Goulmy, Elsa

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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
                                                                                                                                Query Match
Best Local S
Matches 8
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SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                   SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: The HA-1 Antigen FILE REFERENCE: 2183-4285US
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CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                  TYPE: PRT ORGANISM: KIAA0223 partial complementary DNA
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Pred. No. 4.6e+05;
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; LENGTH: 8
; TYPE: PRT
; ORGANISM: HUMAN
US-09-269-250E-38
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-744-549-15
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                                                                 GENERAL INFORMATION:
APPLICANT: Itch, Kyogo et al.
APPLICANT: Itch, Kyogo et al.
TITLE OF INVENTION: HIA-A2-restricted Tumor Antigen Peptides Derived From SART-1
PILE REFERENCE: 0020-4808P
CURRENT PELLICATION NUMBER: US/09/744,549
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: JP H10-212940
PRIOR APPLICATION NUMBER: JP H10-212940
PRIOR FILING DATE: 1998-07-28
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.0
SEQ ID NO 15
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CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
                                                                                                                                                                                                                                                                Sequence 15, Appl
Patent No. 666423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 14
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Best Local (
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APPLICANT: Worley, Paul
APPLICANT: Worley, Paul
APPLICANT: Snyder, Solomon
TITLE OF INVENTION: Huntingtin-associated protein
FILE REFERENCE: 01107.52271
CURRENT APPLICATION NUMBER: US/08/556,419C
CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 25
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APPLICANT: Sharp, Alan
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APPLICANT: Li, Xiao-Jiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 FILE REFERENCE: 58994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Rattus norvegicus
                                                     ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION:
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5; Conserva
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Pred. No.
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3. 4.6e+05;
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GENERAL INFORMATION:

APPLICANT: Vandekerckhove, Joel

APPLICANT: Vandekerckhove, Joel

APPLICANT: Gevaert, Kris

ITILE OF INVENTION: METHODS AND APPARATUS FOR GEL-FREE QUALITATIVE AND

ITILE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AND USES THEREFOR

FILE REFERENCE: VBV-001

CURRENT APPLICATION NUMBER: US/10/394,980

CURRENT PILING DATE: 2003-03-21

PRIOR APPLICATION NUMBER: PCT/EP02/03368

PRIOR APPLICATION NUMBER: US60/278,171

PRIOR APPLICATION NUMBER: US60/278,171

PRIOR APPLICATION NUMBER: US60/318,749

PRIOR APPLICATION NUMBER: US60/318,749

PRIOR APPLICATION NUMBER: US60/318,749

PRIOR APPLICATION NUMBER: US60/318,749

PRIOR PILING DATE: 2001-09-12

PRIOR APPLICATION NUMBER: US60/318,749

PRIOR FILING DATE: 2001-09-12

PRIOR FILING DATE: 2001-09-20

NUMBER OF SEQ ID NOS: 473

COUNTAINS OF SEQ ID NOS: 473
                                                           TITLE OF INVENTION: METHODS AND APPARATUS FOR GEL-FREE QUALITATIVE AND TITLE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AND USES THEREFOR FILE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AND USES THEREFOR FILE OF INVENTION UNDER: US/10/394,980
[CURRENT APPLICATION NUMBER: PCT/FPO2/03368
[PRIOR APPLICATION NUMBER: PCT/FPO2/03368
[PRIOR APPLICATION NUMBER: W560/278,171
[PRIOR APPLICATION NUMBER: US60/278,171
[PRIOR APPLICATION NUMBER: US60/318,749
[PRIOR APPLICATION NUMBER: US60/318,749
[PRIOR APPLICATION NUMBER: US60/318,749
[PRIOR FILING DATE: 2001-09-12
[PRIOR FILING DATE: 200
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US-10-394-980-464
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Best Local Similarity
Thes 5; Conserve
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US-10-394-980-421
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 464, Application US/10394980
Patent No. 6908740
GENERAL INFORMATION:
APPLICANT: Vandekerckhove, Joel
APPLICANT: Gevaert, Kris
TITTE OF TATELY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF T
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SOFTWARE: PatentIn version 3.1 
EQ ID NO 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: MISC FEATURE
OTHER INFORMATION: FR10 - 1624,68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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Local Similarity 71.4%;
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
0; Mismatches 2
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US-08-459-568-20
                                                       RESULT 13
US-08-399-411-20
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                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 12
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9901
TELEPHONE: (619) 535-9949
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
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Best Local Similarity 71.4%;
Matches 5; Conservative
Sequence 20, Application US/08399411
Patent No. 5831008
GENERAL INFORMATION:
                                                                                                                                                                                            Matches
                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 02-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/3:
AFTING DATE: 06-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION: APPLICANT: Huang,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: MISC_FEATURE
OTHER INFORMATION: Table XIX - 1624.68
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPONITION
OF STREM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/459,568
APPLICATION NUMBER: US/08/459,568
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                                                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                              linear
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Pred. No. 4.6e+05;

    Mismatches

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Pred. No. 4.6e+05;
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US-08-516-859A-20
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COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PATENTIA Release #1.0, Version #1.25
SOPTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/399,411
FILING DATE: 06-MAR-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REGISTRATION NUMBER: P-LJ 1264
TELECOMMUNICATION INFORMATION:
TELEPAX: (619) 535-9001
TELEPAX: (619) 535-8949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20, Application US/08516859A
Patent No. 6069231
GENERAL INFORMATION:
APPLICANT: Huang, Shi
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                                                     SOFTWARE: PATENTIN Release #1.0, VE CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/516,859A FILING DATE: 18-AUG-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/399,411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (619) 535-8949 INFORMATION FOR SEQ ID NO:
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TITLE OF INVENTION: Retinoblastoma Protei
TITLE OF INVENTION: Zinc Finger Proteins
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LENGTH: 6 amino acids
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite
                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Retin TITLE OF INVENTION: Zinc NUMBER OF SEQUENCES: 106
FILING DATE: 06-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                        COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release
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STATE: California
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ZIP: 92122
                                                                                                                                                                                                                                                                                                                        CITY: San Diego
STATE: California
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                                                                                                                                                                                                                                                                                                                                                             E: Campbell & Flores LLP
4370 La Jolla Village Drive, Suite 700
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Retinoblastoma Protein - Interacting Zinc Finger Proteins
                                                                                                                                              US/08/516,859A
      US 08/292,683
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Pred. No. 4.6e+05;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 20: US-09-586-472-20
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US-09-586-472-20
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NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1776
TELECOMMUNICATION INFORMATION:
TELEPRAN: (619) 535-9901
TELEPRAN: (619) 535-9949
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20, Application US/09586472
Patent No. 6323335
PATENT INFORMATION:
APPLICANT: Huang, Shi
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
Zinc Finger Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                  TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                            APPLICATION NUMBER: US 09/528,706
FILING DATE: 17-MAR-2000
APPLICATION NUMBER: US 08/516,859
FILING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
APPLICATION NUMBER: US 08/399,411
FILING DATE: 18-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REGERENCE/DOCKET NUMBER: P-LJ 4130
TELECOMMUNICATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 6 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/586,472
FILING DATE: 01-Jun-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
CORDENCE ADDRESSES: Compbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 DDLLE 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: San Diego
STATE: California
COUNTRY: USA
                                                            TYPE: amino acids
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    Mismatches

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Query Match

Guery Match

Guery Match

Best Local Similarity

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Matches 4; Conservative 1; Mismatches 0; Indels 0; Gaps

Qy

4 DLLLE 8

Search completed: April 6, 2006, 16:27:37

Job time: 27.6667 secs
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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq
      length: 0 length: 9
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Match
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37
1 VLXDDLLEA 9
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    Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11B_PUBCOMB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
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US-10-623-176-1
US-10-623-176-2
US-10-623-176-10
US-10-791-217-2
US-10-791-217-2
US-10-861-335-1
US-11-007-740-18
US-11-007-740-20
US-11-007-740-20
US-11-007-740-20
US-11-007-740-20
US-11-007-740-20
US-10-623-176-42
US-10-623-176-45
US-10-623-176-46
US-10-623-176-48
US-10-623-176-48
US-10-623-176-3
US-10-623-176-5
US-10-119-536A-109
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                  Sequence
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20, Appli
41, Appli
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45, Appli
46, Appli
47, Appli
48, Appli
49, Appli
31, Appli
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Query Matches Matches Qy Db	rs-1	CURRENT PRIOR AP PRIOR DI NUMBER O SOFTWARE SEQ ID NO	RESULT 1 US-10-623-: Sequence Sequence GENERAL GENERAL APPLICAL APPLICAL APPLICAL APPLICAL APPLICAL APPLICAL APPLICAL APPLICAL	2 C C C C C C C C C C C C C C C C C C C	28
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imilarity ; Conservat VLXDDLLEA 9 VLXDDLLEA 9	עס	APPLICATION NUMBER: 003- PELICATION NUMBER: 0003- PELICATION NUMBER: 000-01- LING DATE: 2000-01- PELICATION NUMBER: E PELICATION NUMBER: E PELICATION NUMBER: 99-07- PELICATION NUMBER: J P PELICATION NUMBER: J P PELICATION NUMBER: J P P P P	6-1 , Application US/106231 n No. US20040092446A1 FORWARTION: : Goulmy, Els A.J.M. : Hunt, Donald F. : Engelhard, Victor H. INVENTION: HA-1 epitope.		٥
97.3 100. ive	l Sequence Description wherein X co	N NUMBER: 09/ 19 2003-07 VIVIDER: 09/ 2000-01-21 1997-07-23 VIVIDER: EP 1998-07-23 VIVIDER: PT 1998-07-23 VIVIDER: JP 2000-01-24 S: 77 Ver. 2.1	US/106)92446A A.J.M. A.J.M. Uictor Victor	συσωσωσωσωσωσωσωσωσωσωσωσωσωσωσωσωσωσωσ	
; Score 36; DB 4; Len. %; Pred. No. 1.7e+06; 0; Mismatches 0;	nce tion of Artificial Sequence X can be R or H	MBER: US/10/623,176 2003-07-18 ER: 09/489,760 0-01-21 0-01-21 ER: EP 97202303.0 ER: EP CT/NL98/00424 8-07-23 8-07-23 0-01-24 77 7. 2.1	US/10623176 92446A1 A.J.M. d.F. Victor H. Victor H.	-10-275-652-50 -10-952-557-46 -10-952-557-46 -10-956-306-10 -09-833-203-54 -10-057-4758-1 -10-057-4758-1 -10-154-8848-1 -10-623-17-4 -10-623-17-4 -10-623-17-4 -10-919-217-4 -10-919-364-73 -11-007-740-40 -10-166-225A-1 -10-166-225A-1 -10-166-225A-1 -10-166-225A-1 -10-166-225A-1	4 US-10-394-980-464
gth 9; Indels 0; Gaps	: HA-1 peptide	•		100 100 100 100 100 100 100 100 100 100	64
0,				, Appl 1, App 4, App 9, Appl 6, Appl 813, A 813, Appl	ב כ

RESULT 2
US-10-623-176-2
; Sequence 2, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
; APPLICANT: Goulmy, Els A.J.M.
; APPLICANT: Hunt, Donald F.

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PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILLING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILLING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILLING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 10
                                                                                                                   ; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-623-176-10
; Sequence 10, Application US/10623176
; Publication No. US20040092446A1
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                                     Query Match
Best Local Similarity
Matches 8; Conserv
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/489,760 PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/623,176 CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
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TYPE: PRT
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                                                                                                                                                                                                 FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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LOCATION: (1)..(9)
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                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                 TYPE: PRT
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                                       Conservative
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                                       0
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Pred. No. 1.7e+06;
                                                         Score 36; DB 4;
Pred. No. 1.7e+06;
                                           Mismatches
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FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; FEATURE:
, NAME/KEY: MISC_FEATURE
, LOCATION: (3)...(3)
, OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
US-10-791-217-1
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                                                                            , OTHER INFORMATION: obtained from histocompatibility antigen US-10-791-217-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
Best Local Similarity
Matches 8; Conserv
                                        Query Match
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION UNMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION: obtained from histocompatibility antigen
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                                                                                                                   ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                         TYPE: PRT
                                                                                                                                                                                 LENGTH: 9
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    Conservative
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100.0%; Pred. No.
                      97.3%; Score 36; DB 4; 88.9%; Pred. No. 1.7e+06;
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US-10-791-217-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
                                                                                                                                               US-10-861-335-1
                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/10861335
Publication No. US20050031612A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot
TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
CURRENT APPLICATION NUMBER: US/10/861,335
CURRENT FILING DATE: 2004-06-04
CURRENT FILING DATE: 2004-06-04
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Best Local
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LENGTH: 9
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                                                                                       Query Match
Best Local
                                                                        Matches
                                                                                                                                                                                                                                                        SEQ ID NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
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PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
APPLICANT: Engelhard, Victor H
                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR FILING DATE: 2002-12-05
                                                                                                                                                                             LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                              OTHER INFORMATION: HA-1 peptide
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8; Conserv
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VLHDDLLEA
                                    VIXDDLLEA 9
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Pred. No. 1.7e+06;
0; Mismatches 1
                                                                                     Score 36; DB 5;
Pred. No. 1.7e+06;
                                                                      Mismatches
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                                                                                                         Length 9;
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RESULT 8
US-11-007-740-18
; Sequence 18, Application US/11007740
; Publication US/0505033350A1
; GENERAL INFORMATION:
; APPLICANT: GOLLINY, Elsa
; TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
; TREFERENCE: 2799/58994-A
RESULT 10
US-11-007-740-29
; Sequence 29, Application US/11007740
; Publication No. US20050233350A1
; GENERAL INFORMATION:
; APPLICANT: GOULMY, E18a
; TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
; FILE REFERENCE: 2799/58994-A
; CURRENT APPLICATION NUMBER: US/11/007,740
; CURRENT FILING DATE: 2004-12-08
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Sequence 20, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
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Best Local S
Matches 8
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR PILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1 SEQ ID NO 18
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CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Exon
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Pred. No. 1.7e+06;
0; Mismatches 1
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Pred. No. 1.
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1.7e+06;
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; NAME/KEY: MISC FEATURE
; LOCATION: (3). . . (3)
; OTHER INFORMATION: Xaa represents a histidine
US-11-007-740-29
                                                                                                        US-10-623-176-42
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Best Local Similarity
Wetches 7; Conserva
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                                                                                                                            RESULT 12
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
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LENGTH: 9
TYPE: PRT
Sequence 42, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 41, Application US/106 Publication No. US20040092446A1 GENERAL INFORMATION:
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Best Local (
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PRIOR PILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT EPPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
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NAME/KEY: SITE
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9; Conserve
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Pred. No.
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GENERAL INFORMATION:

APPLICANT: Goulmy, Els A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Hunt, Donald F.

APPLICANT: Hunt, Donald F.

APPLICANT: Hagelhard, Victor H.

TITLE OF INVENTION: HA-1 epitopes and uses thereof

FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

FRIOR APPLICATION NUMBER: 09/489,760

PRIOR FILLING DATE: 2000-01-21

PRIOR FILLING DATE: 1997-07-23

PRIOR APPLICATION NUMBER: EP 97202303.0

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR FILING DATE: 1998-07-23

PRIOR APPLICATION NUMBER: DF 97202303.0

PRIOR FILING DATE: 1998-07-23

PRIOR FILING DATE: 1998-07-23

PRIOR FILING DATE: 2000-01-24

NUMBER OF SEQ ID NOS: 77

SEQ ID NO 45

1. FMCTH: 9
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US-10-623-176-45
; Sequence 45, Application US/10623176
; Publication No. US20040092446A1
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-42
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: ECT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: 92000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                             ; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-45
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Best Local
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                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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Local Similarity 87.5%;
es 7; Conservative
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2 LXDDLLEA 9
                                                                       Similarity 7; Conserv
                                                                          Conservative
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                                                                                          86.5%;
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Pred. No. 1.7e+06
                                                                                               Score 32; DB 4;
Pred. No. 1.7e+06;
                                                                     Mismatches
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US-10-623-176-47
; Sequence 47, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
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APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR FILING DATE: 2000-01-21

PRIOR PILING DATE: 1997-07-23

PRIOR APPLICATION NUMBER: BP 97202303.0

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR FILING DATE: 1998-07-23

PRIOR FILING DATE: 2000-01-24

NUMBER OF SEQ ID NOS: 77

SOFTMARE: PatentIn Ver. 2.1
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US-10-623-176-46
US-10-623-176-46
; Sequence 46, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: ECT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
                                                                                                                          NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 47
LENGTH: 9
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Best Local Similarity
Matches 7; Conserv
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
FEATURE:
NAME/KEY: SITE
LOCATION: (1)..(9)
3-10-623-176-46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                        ORGANISM: Artificial Sequence FEATURE:
                                                                                                 TYPE: PRT
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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Search completed: April 6, 2006, 16:50:29
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Post-processing: Minimum Match 0%
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Perfect score:
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Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/USOS NEW PUB.pep:*

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3: /SIDSS/ptodata/2/pubpaa/USO7 NEW PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT NEW PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/USO9 NEW PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*
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37
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Gapop 10.0 , Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	ngth DB	ID	Description
1	36	97.3	9 7	US-11-010-748A-11	Sequence 11, Appl
2	36	97.3	9 7	US-11-010-748A-12	
ω	21	56.8	9	US-10-927-435-63	63
4.	21	56.8	φ 6	US-10-927-634-63	63
5	20	54.1	5 7	US-11-129-143-168	168,
0	20	54.1	5 7	US-11-129-143-169	169,
7	20	54.1	5 7	US-11-129-143-170	170,
80	20	54.1	5 7	US-11-129-143-171	171,
9	20	54.1	5 7	US-11-129-143-172	172
10	20	54.1	5 7	US-11-129-143-173	
11	20	54.1	7 6	US-10-982-440-126	126
12	20	54.1	8 7	US-11-021-305-62	62,
13	20	54.1	8 7	US-11-021-305-65	Sequence 65, Appl
14	20	54.1	8 7	US-11-021-305-66	66,
15	19	51.4	6 7	US-11-207-078-373	
16	19	51.4	7 7	US-11-225-686-2954	Sequence 2954, Ap
17	19	51.4	7 7	US-11-225-686-3877	
18	19	51.4	7 7	US-11-225-686-3914	
19	19	51.4	7 7	US-11-225-686-3915	
20	19	51.4	7 7	US-11-225-686-3916	
21	19	51.4	7 7	US-11-225-686-3932	
22	19	51.4	7 7	US-11-225-686-3941	
23	19	51.4	7 7	US-11-202-009-2954	
24	19	51.4	7 7	US-11-202-009-3877	
25	19	51.4	7	119-11-202-009-3914	

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QY 1 VLXDDILE Db 1 VLHDDILE RESULT 2 VS-11-010-748A-12 VSequence 12, Applica Publication No. US20 GENERAL INFORMATION:	FEATURE: NAME KEY: LOCATION: COTHER INF OTHER INF OTHER INF :-11-010-748 Guery Match Best Local; Matches	PEGINAL PROPERTY OF THE PROPER		44444000000000000000000000000000000000
1 VLX 1 VLH 1 VLH 1 VLH 12, Ap 12, Ap 10n No. INFORMA		11, Applicat: 12, Applicat: 12, Application: 12, Application: 13, Application: 14, Application: 15, Application: 16, Application: 17, Application: 18, Application: 18, Application: 19, Application: 10, Application: 10, Application: 11, Application: 11, Application: 11, Application: 11, Application: 12, Application: 12, Application: 13, Application: 14, Application: 16, Application: 16, Application: 17, Application: 18, Application: 18, Application: 19, Application: 19, Application: 10, Application: 10		119 119 119 119 119 119 118 118 118 118
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	stocompati re 36; DB d. No. 1.4	MET 0,74 5251	ALIGNMENTS	11129-89-11-129-
	patil	METHODS ,, 748A 3251	STYS	
	histocompatibility score 36; DB 7; Le red. No. 1.4e+05; Mismatches 1;		•	US-11-202-009-3916 US-11-202-009-3931 US-11-202-009-3941 US-11-202-009-3941 US-10-93-767A-39 US-10-989-767A-62 US-11-203-137-7 US-11-203-137-7 US-11-203-137-7 US-11-234-786-338 US-11-234-786-338 US-11-129-143-163 US-11-129-143-163
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	antigens ingth 9;	MODULATING		
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RESULT 3
US-10-927-435-63
; Sequence 63, Application US/10927435
; Publication No. US20060045884A1
; GENERAL INFORMATION:
APPLICANT: MOLLDREM, JEFFREY
ITILE OF INVENTION: VACCINES FOR AUTOIMMUNE AND INFECTIOUS DISEASE
; FILE REFERENCE: UTSC:788USC2
; CURRENT FILING DATE: 2004-08-26
; PRIOR APPLICATION NUMBER: 60/489,238
; PRIOR PILING DATE: 2003-08-26
; PRIOR FILING DATE: 2003-08-26
; NUMBER OF SEQ ID NOS: 63
; NUMBER OF SEQ ID NOS: 63
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                     Sequence 63, Application US/10927634
Publication No. US20060045881A1
GENERAL INFORMATION:
APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: ANTI-CANCER VACCINES
FILE REFERENCE: UTSC:788USC1
CURRENT APPLICATION NUMBER: US/10/927,634
CURRENT FILING DATE: 2004-08-26
CURRENT FILING DATE: 2004-08-26
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CURRENT FILING DATE: 2004-12-13
PRIOR APPLICATION NUMBER: PCT/EP03/06251
PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 9
TYPE: PRT
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NAME/KEY: MISC FEATURE
LOCATION: (1)...(9)
OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T
OTHER INFORMATION: ell epitopes
PRIOR APPLICATION NUMBER: 60/489,238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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Pred. No. 1.4e
0; Mismatches
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TYPE: PRT
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TITLE OF INVENTER, ADDRESS, ADDRESS, ADDRESS, ADDRESS, ADDRESS, ADDRESS ADDRES
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Publication No. US20050266518A1

GENERAL INFORMATION:

APPLICANT: BERTY, Alan

APPLICANT: BRETZEL, Werner

APPLICANT: HUMBELLN, Markus

APPLICANT: HUMBELLN, Markus

APPLICANT: HUMBELLN, Markus

APPLICANT: MAYER, Anne F.

APPLICANT: YELISEEV, Alexei A.
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US-11-129-143-168
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SEQ ID NO 168
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TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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NUMBER OF SEQ ID NOS: 63
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TYPE: PRT
ORGANISM: Artificial Sequence
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5; Conserv
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LOPEZ-ULIBARRI,
MAYER, Anne F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BRETZEL, Werner
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Pred. No. 1.4e
1; Mismatches
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Pred. No. 1.4e+05;
2; Mismatches (
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APPLICANT: HUMBELIN, Markus
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C18435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 171
LENGTH: 5
TYPE: PRT
ORGANISM: Bacillus subtilis
US-11-129-143-171
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US-11-129-143-170
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US-11-129-143-169
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                                                                                              Query Match
Best Local &
                                                                            Matches
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Publication No. US20050266518A1
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APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
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4 DDLLE 8
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Pred. No.
                                                                                              Score 20; DB 7; Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
2; Mismatches 0
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                                                                                                                   Length 5;
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APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C3845/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION NUMBER: US/11/129,143
IUMBER OF EGG ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 172
LENGTH: 5
TYPE: PRI
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     RESULT 11
US-10-982-440-126
; Sequence 126, Application US/10982440
; Publication No. US20060018909A1
; GENERAL INFORMATION TAND
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US-11-129-143-172
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US-11-129-143-172
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"---hes 3; Conservative
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; Publication No. US200
; GENERAL INFORMATION:
; APPLICANT: BERRY, Al
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Best Local Similarity 60.0
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APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: WAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1 SEQ ID NO 173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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1 DDILD 5
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Pred. No. 1.4e+05;
2; Mismatches 0
                                                                                                                                                                                                                                                                                            Score 20; DB 7; Lo
Pred. No. 1.4e+05;
2; Mismatches 0;
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RESULT 13
US-11-021-305-65
; Sequence 65, Application US/11021305
; Publication No. US20050282733A1
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US-11-021-305-62
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Best Local Similarity luv.
Conservative
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CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/620,161
PRIOR FILING DATE: 2004-10-19
NUMBER OF SEQ ID NOS: 215
SOFTWARE: Patentin version 3.3
SEQ ID NO 126
LENGTH: 7
LENGTH: 7
                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Prins, Johannes B
APPLICANT: Hutley, Louise J
APPLICANT: Hutley, Louise J
TITLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI169.001CP1
CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR PILING DATE: 2002-06-27
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Matches 4; Conserv
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                                                                     SEQ ID NO 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Graham, Kevin
TITLE OF INVENTION: Angiopoietin-2 Specific Binding Agents
FILE REFERENCE: 04-881-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Prins, Johannes B
APPLICANT: Hutley, Louise J
TITLE OF INVENTION: Differentiation-modulating agents and uses therefor
FILE REFERENCE: DAVI169.001.CP1
CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILLING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR PILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 170
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 170
                                                                                                           NUMBER OF SEQ ID NOS: 170
SOFTWARE: PatentIn version 3.3
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ORGANISM: Homo sapiens
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TYPE: PRT
ORGANISM: mammalian
ORGANISM: mammalian
                                                 TYPE: PRT
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100.0%; Pred. No. 1.4e+05;
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100.0%; Pred. No. 1.4e+05;
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CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.3
SEQ ID NO 66
LENGTH: 8
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US-11-021-305-66
US-11-021-305-66
; Sequence 66, Application US/11021305
; Publication No. US20050282733A1
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US-11-207-078-373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 373, Application US/11207078 Publication No. US20060040307A1 GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Hutley, Louise J
TITLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI169.001CP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: mammalian
COMPUTER READABLE FORN:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/207,078
FILING DATS: 17-Aug-2005
CLASSIFICATION LOTA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/09/721,477
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Local Similarity 100.0%; Pred. No. 1.4e+05;
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ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                           Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
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STATE: California
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
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100.0%; Pred. No. 1.4e+05;
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1: geneseqp1980
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	σ	u	4	ω	2	-	Result No.
26	26	26	26	26	27	27	28	28	28	28	28	28	28	30	31	36	36	37	37	37	45	45	45	Score
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g)				Abu97233 Enzyme pe	Adt02787 Human tum	Abj20115 MHC bindi	Ads81010 Tumour-as	Adq10530 Cercopith	Adk05293 Hepatitis	Adk05291 Hepatitis	Adk68732 Epitope 1	Abg79805 MHC class	Aay10122 T cell ep	Aaw97373 Peptide e	_	Aaw97572 T-cell ep	Aaw99195 Minor his	Adh40334 Human min	Aaw97374 HA-1 R-al	Aaw99197 Minor his	Huma	Aaw97375 HA-1 H-al	Aaw99196 Minor his	Description

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25
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53.3	53.3	53.3		•	53.3	53.3	•	55.6	5	55.6	55.6	55.6	55.6	57.8	57.8	57.8	7.	57.8	57.8	57.8
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Ads78893	Adt39474	Aae28095	Aao21066	Aab70589	Aaw33386	Aar69140	Aar29358	Adc25889	Aae26631	Aab84492	Aag79144	Abu97230	Abu97229	Adu99636	Adt49074	Adt73023	Adm96258	Adm96261	Aae31275	AGM96262
SARS viru	hSARS vir	Human imm	Isomerise	Human imm	Altered I	Endotheli	Endotheli	Yeast GPA	Yeast GPA	Peptide f	Amino ter	Enzyme pe	Enzyme pe	BFA5 tumo	Human BFA	Human RSV	Human ser	Human ser	Human mag	nullidii ser

ALIGNMENTS

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RESULT 1
AAW99196
                                                                                                                                                                                                                                             Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                           20-MAY-1999
                                                                                                                                                                                                                                                                                                         AAW99196;
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                                                                                                                                                                      23-JUL-1997;
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                                                                                                                                                                                                   04-FEB-1999.
                                                                                                                                                                                                                  WO9905174-A1.
                                                                                                                                                                                                                                                                           Minor histocompatibility antigen HA-1 T-cell epitope #2
                                                                                                                                                       (UYLE-) RIJKSUNIV LEIDEN.
                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                      97EP-00202303
                                                                                                                                                                                    98WO-NL000425.
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Goulmy EAJM, Hunt DF, Engelhard VH;

WPI; 1999-153312/13.

A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.

Claim 3; Page 32; 47pp; English.

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auco) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases

Sequence 9 AA;

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RESULT 2
AAW97375
ID AAW9
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  RESULT 3
ADH40333
ID ADH4
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                                                     ADH40333 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
ADH40333;
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-142960/12
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02-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HA-1 H-allele
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Pred. No. 2e+06;
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RESULT 4
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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                                                                                                                                                        Minor histocompatibility antigen HA-1 T-cell epitope #3
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WPI; 1999-142960/12
                   Goulmy E;
                                                           23-JUL-1997;
02-JUN-1998;
                                                                                        23-JUL-1998;
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                                                                                                                                 WO9905313-A2
                                                                                                                                                                        severe aplastic
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                                                           97EP-00202303
98EP-00870125
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RESULT 6
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Matches 8
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                                                                                                                                                           Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                              Strittmatter W,
                                                                                                                                                                                                                                                                                                                                                                                                                                      human; cytostatic; vaccine; SNP profile; minor histocomptibility antigen; mHAg; T
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                                                                                                                   Disclosure; Page 82; 119pp; English.
                                                                                                                                                                                                                    WPI; 2004-082200/08.
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ility antigen; mHAg; T cell epitope.
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The invention relates to a novel method for providing epitopes of allelic variants of antigenic proteins from specific species based on single nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein, identifying, selecting allelic peptide/protein variants, expression product or its fragment encoded by DNA sequence having SNP, creating variant epitopes, selecting epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is

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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                      The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. Pl is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                                                                                                                                                                                                                                          A new minor histocompatibility antigen, diseases and prevent rejection and host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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Sequence 9 AA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-ЛИГ-1998;
                                                                                           treatment of severe aplastic anaemia, leukaemia, and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                and organ transplantation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide;
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Pred. No. 2e+0
0; Mismatches
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2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                          HA-1 - useful to treat immune versus graft disease in bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis neoplastic haematopoietic cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T-cell epitope from the minor histocompatibility antigen
                                                                                                                                                                                                                                                                  Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-versus-host disease.
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                                                                                                                                Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-
                                                                                                                                                                     The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases.
                                                                                                                                                                                                                                             Claim 1; Page 39; 57pp; English
                                                                                                                                                                                                                                                                                                                    WPI; 1999-142855/12.
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                                                                                               Sequence 9 AA;
                                                                                                                       The peptides can also be used and T-cells
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Pred. No. 2e+06;
                                                             Score 36;
Pred. No.
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                                                             DB 2;
2e+06;
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RESULT 9 ADU99853 ID ADU9

ADU99853 standard; peptide; 9 AA.

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Matches 6
                                                                                                                                                                                   Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-MAY-2003;
16-MAY-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention comprises an expression vector that contains a nucleic acid odding a tumor antigen (e.g. BCZ4 or BFY3). The expression vector of invention is useful for the expression of multiple tumor antigens and prevention or treatment of cancer. The present amino acid sequence resents a BCZ4 tumor antigen antigenic peptide that was used in an mple of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression vector comprising a nucleic acid encoding a tumor antigen, . BFA4, BCY1, BFA5, BCZ4, or BFY3, useful for expressing multiple or antigens, or for preventing or treating cancer.
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2003US-0471193P.
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                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                   of HA-1 antigen.
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Pred. No. 2e+0
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2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents an epitope of the minor histocompatibility antigen HA-1. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and
                                                                                                                                                             Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic immunisation; tumour; infectious disease; immunotherapy;
           10-JUL-1997;
                                                                                                                             Synthetic.
                                                                                                                                                   immunisation; tumous
malignant melanoma;
                                                                                                                                                                                                   T cell
                                                                                                                                                                                                                           12-MAY-1999
                                                                                                                                                                                                                                                                       AAY10122 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of genetic aberrances.
                                            10-JUL-1998;
                                                                    21-JAN-1999
                                                                                           WO9902183-A2
                                                                                                                 Human herpesvirus
                                                                                                                                                                                                                                                  AAY10122;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 22; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Typing minor histocompatibility antigen HA-1 - by amplifying identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-142960/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JUL-1997;
02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Goulmy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9905313-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                    _
                                                                                                                                                                                                  epitope/MHC
                                                                                                                                                                                                                                                                                                                                                         1 VLHDDLLEA 9
                                                                                                                                                                                                                                                                                                                                                                                 6
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                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                           (first entry)
           97CA-02209815
97US-00988320
                                            98WO-US014289.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97EP-00202303.
98EP-00870125.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-EP004928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "not specified"
                                                                                                                                                                                                  ligand
                                                                                                                                                    viral disease; hepatitis; AIDS
                                                                                                                                                                                                                                                                                                                                                                                            66.7%;
                                                                                                                                                                                                  SEQ
                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                            Score 30; DB 2;
Pred. No. 2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                   NO:52
                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                        9
                                                                                                                                                               cancer;
                                                                                                                                                                          system;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal; slymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene antigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. malignant melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation that is necessary to keep CTL active, cytotoxic and recirculating through the body. AAY10071 to AAY16339 represent examples of peptide antigens
              WPI; 2002-657506/70
                                                                                                                                                                                                                                                                                   epitope; cytotoxic T lymphocyte response; CTL response; lymphatic system antigen; immunogenic; malignant tumour; carcinoma; melanoma; leukaemia; lymphoma; infectious disease; hepatitis; malaria; measles; tuberculosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 25; 199pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained CTL response, used to treat, e.g. AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kuendig TM,
                                                                                                                                                                                                                                       Cercopithecine
                                                                                                                                                                                                                                                                                                                                   Major histocompatibility complex; MHC; MHC class I molecule; virus;
                                                                                                                                                                                                                                                                                                                                                                   MHC class I molecule, viral epitope #53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABG79805 standard; peptide; 9 AA.
                                                                                                             02-FEB-2001; 2001US-00776232
                                                                                                                                            22-JAN-2002; 2002WO-US002033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunological cytotoxic
                                                                             (CTLI-) CTL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention describes a method of inducing and/or sustaining an
                                              MT,
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5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ø
                                                                                                                                                                                                                                                                      immune deficiency
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IMMUNOTHERAPIES CORP
                                              Simard JJL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                             IMMUNOTHERAPIES CORP
                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first en
                                                                                                                                                                                                                                       herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9
                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62.2%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T lymphocyte (CTL) response
                                                                                                                                                                                                                                                                      syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28;
No.
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2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                      system;
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RESULT 13
ADK68732
ID ADK68
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Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This is particularly useful for treating a mammal having a malignant tumour (e.g. carcinoma, melanoma, leukaemia or lymphoma) or infectious disease (e.g. hepatitis, acquired immune deficiency syndrome (AIDS), malaria, measles or tuberculosis), or in an animal having a predisposition to these diseases. The mammal may be dogs, cats, mice, cattle, sheep, pigs, goats, rabbits, or preferably humans. ABG79753-ABG80319 represent viral epitopes on major histocompatibility complex (MHC) class I molecules, used in the method of the invention. (Updated 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal comprising administering directly to the lymphatic system of the mammal:
(a) an antigen in the form of a polypeptide; (b) a vector comprising a nucleic acid encoding the antigen; or (c) a non-peptide antigen. The method is useful for inducing and/or sustaining CTL response in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Inducing or sustaining immunological cytotoxic T lymphocyte response i mammal, useful for treating a mammal with malignant tumor or infection disease, by directly administering an antigen to the lymphatic system
                                                                                                                                                (SIMA/)
(DIAM/)
(QIUZ/)
(LEIX/)
                                                                                                                                                                                                                                                                                                                                                                                       protozoacide; fungicide; T-cell activator; vaccine; housekeeping epitope; cytotoxic T lymphocyte; CTL; adoptive immunotherapy; neoplastic cell;
                                                                                                                                                                                                                                                                                                                                                                                                                        epitope liberation; substrate; proteasome; cytostatic; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADK68732 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 19; 73pp;
                                                                                                                  Simard JJL,
                                                                                                                                                                                                                          07-NOV-2001; 2001US-0336968P
                                                                                                                                                                                                                                                      07-NOV-2002; 2002US-00292413.
                                                                                                                                                                                                                                                                                                                                             Human herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Epitope liberation-related peptide SeqID95.
                                                                                        WPI; 2004-167209/16.
                                                                                                                                                                                                                                                                                                                 US2003228634-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity es 5; Conserv
                                                                                                                                                ) DIAMOND D (
) QIU Z.
) LEI X.
                                                                                                                                                                                                                                                                                                                                                                           bacterium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ALHDDI 6
                                                                                                                                                                            SIMARD J J L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A
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                                                                                                                     Diamond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                           protozoan; fungus; housekeeping proteasome system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             62.2%;
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Pred. No.
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Disclosure; SEQ ID NO 95; 67pp; English

inversion by contacting substrate polypeptide comprising epitope interest, with proteasome, and assaying for liberation of epitope. Identifying polypeptide suitable for epitope e.g., housekeeping epitope, liberation by contacting substrate polypeptide comprising epitope of

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RESULT 14
ADKOS291
ID ADKOS
XX ADKOS
XX ADKOS
XX PATHO
DT 06-MP
XX PATHO
CKW PATHO
CKW PATHO
CKW HIA---
CXX Hepat
XX WO200
XX WO200
XX WO201
XX WO1;
XX WAT;
XX New
PT Vien
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Best Local :
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                                                                                                                preventing
                                                                                                                                                                                                                                                                                                                                                    Mattner F, Schmidt W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-JUL-2002;
11-JUL-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HLA-allele; CTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pathogenic virus; alternative reading frame; antigenic determinant; virucide; vaccine; therapeutic agent; infection; epitope peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus CTL epitope peptide #3121.
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                                                                                                            polypeptide encoded by an alternative reading us comprising an antigenic determinant, useful venting an infection with the pathogenic virus.
                                                                                                                                                                                                                                                                        2004-169243/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4
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5; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002AT-00001124.
2003EP-00450171.
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Pred. No. 2e+0
1; Mismatches
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                                                                                                                                                        frame of a pathogenic for treating or
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                                                                                                                                                    treating
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Claim 14; Page 114; 220pp; English

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Length

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RESULT 15
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Best Local S
Matches 5
                                           This invention relates to a novel polypeptide encoded by an alternative reading frame of a pathogenic virus, where the polypeptide starts with a methionine amino acid residue, which comprises an antigenic determinant and more than 7 amino acid residues. The invention may be useful for the production of compounds with a virucide activity or the development of a vaccine. The polypeptide or its fragments may be useful as a therapeutic agent. It is also useful for the manufacture of a medicament for treating or preventing an infection with the pathogenic virus. The present sequence is that of a hepatitis C virus CTL epitope peptide of the
Sequence 9
                                                                                                                                                                                                                                                 New polypeptide encoded by an alternative reading frame of a pathogenic virus comprising an antigenic determinant, useful for treating or preventing an infection with the pathogenic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pathogenic virus; alternative reading frame; antigenic determinant;
virucide; vaccine; therapeutic agent; infection; epitope peptide;
HLA-allele; CTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a novel polypeptide encoded by an alternative reading frame of a pathogenic virus, where the polypeptide starts with a methionine amino acid residue, which comprises an antigenic determinant and more than 7 amino acid residues. The invention may be useful for the production of compounds with a virucide activity or the development of a vaccine. The polypeptide or its fragments may be useful as a therapeutic agent. It is also useful for the manufacture of a medicament for treating
                                                                                                                                                                                                                  Claim 14; Page 114; 220pp; English.
                                                                                                                                                                                                                                                                                                                    WPI; 2004-169243/16.
                                                                                                                                                                                                                                                                                                                                                     Mattner F,
                                                                                                                                                                                                                                                                                                                                                                                                                        24-JUL-2002; 2002AT-00001124.
11-JUL-2003; 2003EP-00450171.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus.
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Search completed: April 6, 2006, 16:19:13

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hypothetical protein protein kinase C i elastase - Pseudom urine glycopeptide transaldolase (EC venom protein HR-3 macrophage cytotox trypsin (EC 3.4.21 T-cell receptor be T-cell receptor be cerebellar degener T-cell receptor be cerebellar degener C-cell receptor be acetylcholinestera calsequestrin, fas T-cell receptor be Sexotoxin A - Strep chlorophyll a/b-bi
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phosphoenolpyruvat
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hucolin, 75K chain
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RESULT PQ0663

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3 IIHKNL 8

QY 1 VLHDDL 6	Query Match 37.8%; Score 17; DB 2; Le Best Local Similarity 33.3%; Pred. No. 2.8e+05; Matches 2; Conservative 3; Mismatches 1;	RESULT 2 \$55696 \$55696 C;Species: Trypanosoma brucei C;Species: Trypanosoma brucei C;Date: 28-Oct-1995 #sequence_revision 03-Nov-1995 #tex C;Accession: \$55696 R;Hunt, M.; Koehler, P. Blochim. Biophys. Acta 1249, 15-22, 1995 A;Title: Purification and characterization of phosphoen A;Reference number: \$55696; MUID:95284106; PMID:7766679 A;Accession: \$55696 A;Status: preliminary A;Molecule type: protein A;Residues: 1-9 <hun> A;Cross-references: UNIPROT:Q7M3S5; UNIPARC:UPI000017B5</hun>	Oy 2 LHDD 5 : : Db 3 MHDE 6	Query Match 40.0%; Score 18; DB 2; Best Local Similarity 50.0%; Pred. No. 2.8e+05; Matches 2; Conservative 2; Mismatches 0	RESULT 1 146868 alpha-myosin heavy chain - rabbit (fragment) C;Species: Oryctolagus cuniculus (domestic rabbit) C;Species: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_c C;Accession: 146868 C;Accession: 146868 R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984 A;Title: Characterization of genomic clones specifying rab A;Reference number: 146868; MUID:84221901; PMID:6328491 A;Accession: 146868 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-7 <fri> A;Cross-references: UNIPROT:Q28742; UNIPARC:UPI0000087938;</fri>	ALIGNMENTS	30 12 26.7 9 2 S65913 31 12 26.7 9 2 PH0108 32 12 26.7 9 2 PH0562 33 12 26.7 9 2 B30572 34 11 24.4 5 2 T10994 36 11 24.4 6 2 T1179 37 11 24.4 7 2 S25266 39 11 24.4 7 2 S25266 39 11 24.4 8 2 S22428 40 11 24.4 8 2 S22428 41 12 24.4 8 2 S26915 42 11 24.4 9 2 S66419 43 11 24.4 9 2 PT0272 44 11 24.4 9 2 PT0272 45 11 24.4 9 2 PT0272
	Length 9; ; Indels 0; Gaps 0;	soma brucei fov-1995 #text_change 09-Jul-2004 of phosphoenolpyruvate carboxykinase from Tr: pMID:7766679		Length 7; ; Indels 0; Gaps 0;	#text_change 09-Jul-2004 ing rabbit alpha- and beta-ventricul 8491 087938; GB:K01698; NID:g165538; PIDN		pyrimidine synthes late G1-69 protein T-cell receptor be T-cell receptor be copper resistance hypothetical prote phosphoglycerate t pilE protein - Esc Ig heavy chain CRD chitin-binding pro 158K exoantigen - ferredoxin a2 - Ja tetrameric protein Ig heavy chain CRD transaldolase (EC octamer-binding pr

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membrane protein - porcine epidemic diarrhea virus (isolate Belgian CV777) (fragment) C;Species: porcine epidemic diarrhea virus C;Date: 14-Jul-1994 #sequence_revision 14-Jul-1994 #text_change 08-Oct-1999 C;Accession: PQ0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPARC:UPI0000170FAC; GB:Z14976; C;Comment: This virus is coronavirus related to human C;Keywords: membrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A, Title: Sequence determination of the nucleocapsid protein issible gastroenteritis virus.
A, Reference number: JQ2191; MUID:93389433; PMID:8397280
A, Accession: PQ0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Bridgen, A.; Duarte, M.; Tobler, K.; Laude, H.; Ackermann, J. Gen. Virol. 74, 1795-1804, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-7 < BRI>
                                                                                                                                                        hypothetical protein 8 [imported] - Pseudomonas aeruginosa (fragment) C;Species: Pseudomonas aeruginosa
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 31-Dec C;Accession: Pc4131
R;Kawasaki, S; Arai, H.; Igarashi, Y.; Kodama, T.
Gene 167, 87-91, 1995
A;Title: Sequencing and characterization of the downstream region of ty for biosynthesis of heme d1.
A;Reference number: JC4552; MUID:96144254; PMID:8566817
A;Accession: PC4131
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A,Molecule revo. Nun
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C;Species: Homo sapiens (man)
C;Date: 15-Feb-1997 #sequence_revision
C;Accession: S68004
                                                                                                                                                                                                                                                                                                                                                               RESULT 5
PC4131
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A;Status: preliminary
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                                                                        A;Cross-references: UNIPROT:P95412; UNIPARC:UPI000017A96A; A;Note: this ORF is not annotated in GenBank entry PSENIRC; C;Superfamily: Pseudomonas stutzeri nirD protein
                                                                                                                              A; Molecule type: DNA
A; Residues: 1-8 < KAW>
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 Query Match

Best Local Similarity 100.0%; I

Matches 3; Conservative 0;
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         Score 16; DB; Pred. No. 2.8
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Pred. No.
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3. 2.8e+05;
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5. 2.8e+05;
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     DB 2;
3. 2.8e+05;
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                                            Length 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein kinase C inhibitor - rat (fragment)
C;Species: Rattus norvegicus
C;Date: 25-Aug-2003 #sequence_revision 25-Aug-2003 #text_change 25-Aug-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: prot
A; Residues: 1-6 < GRA>
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S20446
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C;Species: Pseudomonas aeruginosa
C;Date: 19-Mar_1997 #sequence_revision 24-Jul-1997 #text_change
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to the Protein Sequence Database, June 2003 A;Description: Purification of PKCI from rat liver.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Accession:
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                                                                                                      R;Kessler, B.; Safrin, M.; Peretz, M.; Burstein, Y. FEBS Lett. 299, 291-293, 1992
A;Title: Identification of cleavage sites involved in A;Reference number: S20446; MUID:92183956; PMID:154450
A;Accession: S20446
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A; Residues: 1-7 < NEG>
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                                   A; Cross-references:
                                                    A; Molecule type: protein A; Residues: 1-7 < KES>
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                                                                                       A;Status: preliminary
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Best Local S
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     31.1%; Score 14;
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pred. No. 2.8e+05;
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2.8e+05;
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Pseudom

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venom protein HR-3 - oriental hornet (fragment)
('SSpecies: Vespa orientalis (oriental hornet)
('SDecies: Vespa orientalis (oriental hornet)
('Date: 29-Jan-1993 #sequence_revision 29-Jan-1993 #text_change 09-Jul-2004
('Accession: S10920
R;Tuichibaev, M.U.; Akhmedova, N.U.; Kazakov, I.; Korneev, A.S.; Gagel'gans
Blochemistry (N.Y.) 53, 183-190, 1988
Blochemistry (N.Y.) 53, 183-190, 1988
A;Title: Low-molecular-weight peptides of venom of the giant hornet Vespa o
A;Reference number: S06445
A;Accession: S10920
                                                                                                                                                                                                  RESULT 11
S10920
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R;Sun, S.C.; Joris, L.; Tsolas, O.
Arch. Biochem. Biophys. 178, 69-78, 1977
A;Title: Purification and crystallization of transaldolase isozyme I and A;Reference number: A12872; MUID:77110646; PMID:556924
A;Accession: A12872
A;Molecule type: protein
A;Residues: 1-9 cSUN-
A;Residues: 1-9 cSUN-
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C;Date: 20-Jun-2000 #sequence_revision 20-Jun-2000 #text_change 16-Aug-2004
C;Accession: A03188
R;Lote, C;J; Weiss, J.B.
Biochem. J. 123, 25P, 1971
A;Title: Identification in urine of a low-molecular-weight polar glycopeptide containing A;Reference number: A03188; MUID:72062338; PMID:5126885
A;Accession: A03188
A;Molecule type: protein
A;Residues: 1-8 <LOT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transaldolase (EC 2.2.1.2) I - yeast (Pichia jadinii) (fragment)
C;Species: Pichia jadinii, Candida utilis
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:P02729; UNIPARC:UPI000012B849
C;Comment: The identity of the glycoprotein from which this peptide is re has also been found (see PIR:XGHUE).
C;Keywords: glycoprotein
F;1/Binding site: carbohydrate (Cys) (covalent) #status experimental
A;Molecule type: protein
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100.0%; Pred. No. 2.8e+05;
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2; Mismatches 0;
                                                                                             Korneev, A.S.; Gagel'gans,
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                                                       giant hornet Vespa orientalis.
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T-cell receptor beta chain V-D-J region (clone 13) (Species: Rattus norvegicus (Norway rat) (C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 C;Accession: PH0942 R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vande
                                                                                                                                                                                                                                                                                                                                  밁
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C;Keywords: hydrolase; protein digestion; serine proteinase;
F;1-8/Domain: activation peptide #status experimental <APT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    trypsin (EC 3.4.21.4) precursor - minke whale (fragment) c;Species: Balaenoptera acutorostrata (minke whale, lesser rorqual) C;Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004 C;Accession: A61328
                                                                                                                                                                                                                                                                                                                                                                                               Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: prote
A; Residues: 1-8 < BRI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biochim. Biophys. Acta 386, 244-255, 1975 A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, Balaeno, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual, A; Title: N-terminal amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the lesser rorqual amino acid sequence of trypsinogen from the 
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Exp. Hematol. 19, 704-709, 1991
A;Title: Purification and amino acid analysis of a human macrophage cytotoxicity-induci A;Reference number: A60427; MUID:91372335; PMID:1909970
A;Accession: A60427
                                                                                                                                                                                    PH0942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Bricteux-Gregoire, S.; Schyns, R.; Florkin, M.; Emmens, M.; Welling, G.W.; Beintema, Biochim. Biophys. Acta 386, 244-255, 1975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
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C;Species: Homo sapiens (man)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 12
A60427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Reference number: A61328; MUID:75146765; PMID:1125273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: protein A; Residues: 1-9 < JON>
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Local Similarity 28.6%;
                                                                                                                                                                                                                                                                                                                                  3 IDDD 6
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3; Conserv
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Pred. No.
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Pred. No. 2.8e+05;
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Pred. No. 2.8e+05;
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
Vandenbark,
                                                                   #text_change 30-May-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 8;
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J. Exp. Med. 174, 1467-1476, 1991
A, Title: Analysis of T cell receptor beta chains in Lewis rats with experimental allergi A, Reference number: PHO891; MUID:92078857; PMID:1836012
A, Accession: PHO942
A, Molecule type: mRNA
A, Residues: 1-9 <GOL>
A, Cross-references: UNIPARC:UDI000017C9DD
A, Experimental source: complete Freund's adjuvant-immunized lymph node
A, Mote: the authors translated the codon TGC for residue 2 as Ala
C; Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 15
140697
biotin A - Citrobacter freundii (fragment)
C.Species: Citrobacter freundii
C.Species: Citrobacter freundii
C.Species: Citrobacter freundii
C.Pate: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C.Pate: 12-Aug-1996 #sequence_revision 12-Aug-1996 #seque
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Search completed: April Job time: 19.6667 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 26.7%; Score 12; DB 2; Length 4; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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5 LLE 7
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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    92:
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                                                                                                                                                                                                                                                                        Score
                                                                                                                                                                                                                                                                                                                                                                                                                  seq length: 0
seq length: 9
    Query
Match
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Gapop 10.0 ,
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1: uniprot_sprot:*
2: uniprot_trembl:*
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   US-10-791-217A-2
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Q47556 ECOL1
Q28742 RABIT
Q28742 RABIT
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Q99182 9SMEG
Q726GO HUMAN
FAR8 MACRS
Q70SM2 HUMAN
Q4X981 PLACH
Q8LPTS MAIZE
P82568 STRPY
Q15900 HUMAN
         TAL1 PĪCJA
Q7M4R5 HUMAN
Q7M471 VESOR
Q9FSZ2 CICAR
Q9FSZ2 CICAR
Q8CG39 RAT
Q9QZA8 MOUSE
Q6Q7G0 RANRI
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Q85710 9RETR
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Q80TD7-9HIVI
Q66205 9CORO
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Q9XJN0_9VIRU
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GLUR_HUMAN
Q9BFA7_MACPR
P72279_RHOGO
Q93SR0_STAEP
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                                                                                                                                                  Q778x5 plasmodium
Q47556 escherichia
Q28742 oryctolagus
Q7m3s5 trypanosoma
Q99182 gnatholebia
Q72690 homo sapien
P83281 macrobrachi
Q70sm2 homo sapien
Q4x981 plasmodium
Q81pt5 zea mays (m
P82568 streptococc
Q15900 homo sapien
Q4x127 plasmodium
 Q9bfa7
P72279
Q93e79
Q93e740
P17440
Q7m4r5
Q7m471
Q9f2a3
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4 pseudomonas

9 homo sapien

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18 pagergillus

19 homo sapien
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8 streptococc
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RESULT 2
Q47556_ECOLI PRELIMINARY;
ID Q47556,
AC Q47556;
DT 01-NOV-1996 (TrEMBLrel. 01, C:
DT 01-NOV-1998 (TrEMBLrel. 08, L:
DT 01-NOV-1998 (TrEMBLrel. 08, L:
GN Name=pyrI;

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regulatory chain (Fragment).

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o o	vati	data 528; otein ; 10	re 419:512-519(2002). CAUTION: The sequence EMBL/GenBank/DDBJ who	sequence and comparative Plasmodium yoelii yoeli	ungham D.A., Preiser P.R., Lin L.H., Janse C.J., Wate Derg S.L., Venter J.C., Fr	S.J., van Aken S.E., Riedmuller., Quackenbush J., Sedegah M., S.L., Yates J.R. III, Raine J.D.,	Ermolaeva N	55706; PubMed=12368865;	ence 9; oraca;	yoelii yoelii	(TrEMBLrel. (TrEMBLrel. (TrEMBLrel. protein.	PREL IMINARY		99	99	999	99	. .	ω ω ·	φ φ
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DT Alpha-my
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RESULT 4
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Bacteria; Proteobacteria; Gammaproteobacteria;
Enterobacteriaceae; Escherichia.
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Proc. Natl. Acad. S
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MEDLINE=82275057; PubMed=7051000;
Pauza C.D., Karels M.J., Navre M.
"Genes encoding Escherichia coli descriptions of the coli descriptions of the coli descriptions of the coli descriptions of the coli
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Alpha-myosin heavy chain (Fragment).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Lagomorpha; Leporida
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Hoover T.A., Roof W.D., Foltermann
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NUCLEOTIDE SEQUENCE:
MEDLINE=84221901; PubMed=6328491;
Umeda P.K., Sinha A.M., Hsu H.J.,
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Q28742;
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Proc. Natl. Acad. Sci. U.S.A. 81:3044-3048(1984).
EMBL; K01698; AAA31415.1; -; Genomic_DNA.
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Q7Z6G0 HUMAN PRELIMINARY; P
Q7Z6G0;
01-OCT-2003 (TrEMBLrel. 25, Crea
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Funarate hydratase (Fragment).
Homo sapiens (Human).
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099182;
01-MAY-1999 ('
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Phosphoenolpyruvate carrboxykinase.
Trypanosoma brucei.
Eukaryota: Euglenozoa; Kinetoplastida; Trypa
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SEQUENCE 7 AP
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Cytochrome oxidase I (Fragment).
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**CLEOTIDE SEQUENCE.

**RA Wei M.-H., Nickerson M.L., Toro J.R.;

RL Submitted (MAY-2003) to the EMBL/GenBank/DDBJ

REMBL; AY299638; AAP57532.1; -; Genomic_DNA.

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T NON_TER 8 8

P SEQUENCE 8 AA.
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28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
95-JUL-2004 (Rel. 44, Last annotation update)
FMRFamide-like neuropeptide FLPB (VSHNNFLRF-amide).
Macrobrachium rosenbergii (Giant fresh watter prawn).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Caridea
Palaemonoidea; Palaemonidae; Macrobrachium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=21107394; PubMed=11179812; DOI=10.1016/S0196-9781(00)00382-X; Sithigorngul P., Saraithongkum W., Longyant S., Panchan N., Sithigorngul W., Petsom A., Sithigorngul W., Petsom B., Three more novel FWRFamide-like neuropeptide sequences from the eyestalk of the giant freshwater prawn Macrobrachium rosenbergii."; Peptides 22:191-197(2001).
                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                            Amidation; Direct MOD_RES 9
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=79674;
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les 3; Conserv
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MASS SPECTROMETRY: MW=1133.8; METHOD=MALDI; RANGE=1-9; NOTE=Ref.1.

SIMILARITY: Belongs to the FARP (FWRFamide related peptide)
                                                                                                                                                                                                                                                                                                                                                                                                                                          GO:0007218; P:neuropeptide signaling pathway; IDA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 s Swiss-Prot entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its as long as its content is in no way modified and this statement is not
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                             PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                  protein sequencing;
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e EMBL/GenBank/DDBJ databases.
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Pred. No. 2.2e+06;
2; Mismatches 2;
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RESULT Q8LPT5 ID QE

MAIZE

QBLPT5_MAIZE PRELIMINARY;

PRT;

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RESULT 9
Q4X981 PLACH
ID Q4X981;
D7 13-SEP-2
D8 Hypothet
GN NCHITAX
RN [1]
RA Berriman
RA James K.
RA Quail M.,
RA Bidwell:
RA Gunil M.,
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                                                                                                                                   Matches
                                                                                                                                                                   Query Match
Best Local
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Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.K.,
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C.,
Janse C.J., Barrell B., Turner C.M.E., Waters A.P., Sinden R.S.;
"A comprehensive survey of the Plasmodium life cycle by genomic,
transcriptomic, and proteomic analyses.",
Science 307:82-86 (2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plasmodium chabaudi.
Eukaryota; Alveolata; Apicomplexa;
NCBI_TaxID=5825;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical protein (FORFNames=PC404684.00.0;
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13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                          Hypothetical protein.
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Q4X981;
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Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                    preliminary data.
; CAAJ01008648; CAH86543.1;
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3 DTMEA 7
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PRESULT 11

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DE HOMO SAP
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                                                                                             Q15900 HUMAN PRELIMINARY;
Q15900;
01-NOV-1996 (TrEMBLrel. 01, Cr
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01-DEC-2001 (TrEMBLrel. 19, La
01-DEC-2001 (Clone XP7B11A)
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Submitted (APR-2002) to the EMBL/GenBan Submitted (APR-2002) to the EMBL/GenBan Submit AN
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Coolbaugh M.I., Chinault C.A., Baldini
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                       Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A., Berriman M., Florens L., Janssen C.S., Pain A., Christophides G. Barriman M., Florens L., Harris B., Harris D., Churcher C., James K., Rutherford K., Harris B., Harris D., Churcher C., Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J., Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S., "A comprehensive survey of the Plasmodium life cycle by genomic,"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Caskey
                                                                                                                                                                                                                                                                                                                                                                                                                         Plasmodium chabaudi.
Bukaryota; Alveolata; Apicomplexa;
                                                                                                                                                                                                                                       preliminary data.
EMBL; CAAJ01003387; CAH79935.1; -; Genomic_DNA.
                      01-NOV-1999 (TrEMBLrel.
01-NOV-1999 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                   transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                                                        Q9XJNO_9VIRU
Q9XJNO;
 Bacteriophage phi-10.
            P10 (Fragment)
                                                                                                                                                                                                                                                           CAUTION: The sequence shown here is derived EMBL/GenBank/DDBJ whole genome shotgun (WGS)
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Last annotation update)
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pred. No. 2.2e+06;
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Last sequence update)
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Pred. No.
                                                                                                                                                                                    Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Haemosporida; Plasmodium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=9350412; PubMed=10419946;
Mindich L., Qiao X., Qiao J., Onodera S., Romantschuk M.,
Hoogstraten D.;
"Isolation of additional bacteriophages with genomes of segmented
double-stranded RNA.";
J. Bacteriol. 181:4505-4508(1999).
EMBL; AR125675; AAD225555.1; -; Genomic_RNA.
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                                                                                                                                                                                                                                          Tryptophan biosynthesis. NON_TER 6 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                  InterPro; IPR000847; HTH Lysr.

PROSITE; PS50931; HTH LYSR; PARTIAL.

Activator; Amino-acid biosynthesis; Aromatic amino acid biosynthesis; DNA-binding; Transcription; Transcription regulation;
                                                                                                                                                                                                                                                                                                                                                                                             EMBL; X13299; CAA31660.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-89335826; PubMed-2503057; DOI=10.1016/0300-9084(89)90183-1; Eberly L., Crawford I.P.; "DNA sequence of the tryptophan synthase genes of Pseudomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pseudomonas putida.
Bacteria; Proteobacteria; Gamm
Pseudomonadaceae; Pseudomonas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; dsRNA viruses; NCBI_TaxID=90889; [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                               removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biochimie 71:521-531(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-PPG1
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nes 2; Conserv
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SIMILARITY: Contains 1 HTH lysR-type DNA-binding domain.
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US-09-489-760-2
US-09-269-250B-18
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US-09-269-250B-38
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US-09-269-250B-38
US-09-25-819-5
US-09-803-951-5
US-09-803-951-5
US-09-803-951-7
US-08-98-923A-7
US-08-98-923A-7
US-08-169-323B-8
US-08-169-323B-8
US-08-159-339A-370
US-08-189-339A-370
US-08-189-339A-370
US-08-208-036-7
US-08-208-036-7
US-08-208-036-7
US-08-208-036-9
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US-08-208-036-9
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US-08-268-419-14
US-09-173-941-82
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Biocceleration
Sequence 38, Appli
Sequence 5, Appli
Sequence 5, Appli
Sequence 7, Appli
Sequence 4, Appli
Sequence 8, Appli
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Sequence 30, Appli
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Sequence 14, Appli
Sequence 18, Appli
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Sequence
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2, Appli
18, Appl
5, Appli
5, Appli
1, Appli
1, Appli
4, Appli
38, Appl
                                                                          Sequence 2, Application US/09489760
Patent No. 6678375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1988-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
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US-09-489-760-2
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       ORGANISM: Histocompatibility antigen
                                     TYPE:
                                                       LENGTH:
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21	21	21	21	21	21	21	21	21	21	21	21	21	21	21	21	22	22
46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	46.7	48.9	48.9
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US-08-459-568-78	US-08-459-568-72	US-08-343-602-2	US-07-671-757-30	US-09-528-706-75	US-09-586-472-75	US-09-419-826-37	US-09-419-826-1	US-08-516-859A-75	US-08-399-411-75	US-08-459-568-75	US-09-528-706-20	US-09-586-472-20	US-08-516-859A-20	US-08-399-411-20	US-08-459-568-20	US-08-873-235B-11	US-09-217-609A-11
Sequence 78, Appl	Sequence 72, Appl	Sequence 2, Appli	Sequence 30, Appl	Sequence 75, Appl	•	Sequence 37, Appl	Sequence 1, Appli	•	-	Sequence 75, Appl	•	•	Sequence 20, Appl	Sequence 20, Appl	Sequence 20, Appl	•	Sequence 11, Appl

ALIGNMENTS

Sequence 20, Application US/09269250E Patent No. 6830883 GENERAL INFORMATION: APPLICANT: GOULNY, Elsa TITLE OF INVENTION: METHOD FOR TYPING OF MING FILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: US/09/269,250E CURRENT FILING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 20 LENGTH: 9 US-09-269-250E-20 US-09-269-250E-20 Query Match Best Local Similarity TYPE: PRT ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Exon 1 VLHDDLLEA 1 VLHDDLLEA 9 9; 100.0%; nilarity 100.0%; Conservative (9 0; Score 45; DB 2; Pred. No. 4.6e+05; Mismatches OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 Length 9; Indels Fragments <u>.</u> Gaps 0,

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Best Local Similarity
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US-09-269-250E-18
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US-09-489-760-5
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                                                                                                                                                                                                                                                             APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hout, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER RILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
RESULT 5 '
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Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1
SEQ ID NO 18
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                            Matches
                                                                                                                        Query Match
Best Local
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                                                                                                                                                                                       ORGANISM: KIAA0223 partial complementary DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                       TYPE: PRT
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
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US-09-489-760-4
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PATERIX NO. 683083
PATERIX INFORMATION:
APPLICANT: GOUND, E18a
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATERIX VERSION 3.1
SEQ ID NO 29
LENGTH: 9
                 RESULT 7
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Best Local S
Matches
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                                                                                                                                    Matches
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Best Local
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CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 198-07-23
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                    LENGTH: 9
TYPE: PRT
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LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine (H) or an arginine
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ORGANISM: Human
FEATURE:
                                                                                                                                                                                                                                                                ORGANISM: Histocompatibility antigen FEATURE: NAME/KEY: PEPTIDE
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OTHER INFORMATION:
OTHER INFORMATION:
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Pred. No. 4.6e+05;
0; Mismatches 1
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Pred. No.
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Sequence 4, Application US/09489760
Patent No. 6878375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin

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US-09-269-250E-38
; Sequence 38, Application US/09269250E
; Sequence 38, Application US/09269250E
; Patent No. 6830883
; GENERAL INFORMATION:
; APPLICANT: Goulmy, Elsa
; TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
; FILE REFERENCE: 58994
; CURRENT APPLICATION NUMBER: US/09/269,250E
; CURRENT FILING DATE: 1999-05-21
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , LOCATION: (2)..(7)
; OTHER INFORMATION: AMINO ACIDS X REPRESENT LEUCINE OR ISOLEUCINE
; OTHER INFORMATION: RESIDUES
US-09-489-760-4
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Best Local Similarity
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Patent No. 6225097
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Best Local Similarity
Matches 6; Conserv
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                                                   GENERAL INFORMATION:
APPLICANT: Obata, Shusei
APPLICANT: Nishino, Tokuzo
APPLICANT: Koyama, Tanetoshi
APPLICANT: Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: HUMAN
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TYPE PRT
TYPE (AND TYPE)
ORGANISM: Histocompatibility antigen
FEATURE:
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                                   CORRESPONDENCE ADDRESS
   ADDRESSEE:
STREET: 15
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8: KENYON & KENYON
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88.9%;
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Pred. No. 4.6e+05;
0; Mismatches 0;
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Pred. No.
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RESULT 10
US-09-808-126-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Application Patent No. 6410280 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 4; Conserv
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FILING DATE: 19-FEB-1998
CLASSIFICATION: 435
FRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 251675
FILING DATE: 17-SEP-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                    ZIP: 20005

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/808,126
FILING DATE: 08-May-2001
CLASSIFICATION UNMBER: US/09/808,126
FILING DATE: OWNERS: 09/025,819
PRIOR APPLICATION UNMER: 09/025,819
PRIOR APPLICATION UNMER: 09/025,819
APPLICATION UNMER: 09/025,819
APPLICATION INFORMATION:

NEW YORK OF THE PRIOR 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: 202-220-4200
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REGISTRATION NUMBER: 39,546
REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Obata, Shusei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 УТНОО 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 1500 K Street, N.W. CITY: Washington
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: KENYON & KENYON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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Koyama, Tanetoshi
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Pred. No. 4.6e+05;
1; Mismatches 0
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RESULT 11
US-09-803-951-5
; Sequence 5, Application
; Patent No. 6413761
; GENERAL INFORMATION:
                                                                    ; TOPOLECULE TYPE: POPTION: SEQ ID NO: 5: US-09-803-951-5
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                                                                                                                                                                                                         APPLICATION NUMBER: 09/025,819
FILING DATE: <UNKNOWN-
ATTORNEY/AGERT INFORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
REFERENCE/DOCKET NUMBER: 10235/2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-220-4200
TELEPHONE: 202-220-4201
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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Matches 4; Conservative
Ousry Match
Best Local Similarity 80.0
Matches' 4; Conservative
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/803,951
FILING DATE: 13-Mar-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION: DATA:
OF THE PRIOR OF THE PARTY OF THE PRIOR APPLICATION DATA:
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MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 5:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-220-4200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: KENYON & KENYON
STREET: 1500 K Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nishino, Tokuzo
Koyama, Tanetoshi
Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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STRANDEDNESS: <Unknown>
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                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: DC
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                                                                                                                                                                                                 LENGTH:
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                                                                                                                                                                                               7 amino acids
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80.0%;
                    55.6%;
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    Score 25; DB 2; Length 7; Pred. No. 4.6e+05; Indels 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                    10235/2
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US-09-305-923A-7
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                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Ostanin, Kirill
APPLICANT: Silverman, Lauren
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TITLE OF INVENTION: YEAST CELLS HAVING MUTATIONS IN Stp22 AND USES THEREFOR FILE REFERENCE: CPI-091
CURRENT APPLICATION NUMBER: US/09/305,923A
CURRENT FILING DATE: 1999-05-05
PRIOR APPLICATION NUMBER: 60/084,420
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ABCII (text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/582,333A
FILING DATE: 17-JAN-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Catherine J. Kara
REGISTRATION NUMBER: 41,106
REFERENCE/DOCKET NUMBER: 41,106
REFERENCE/DOCKET NUMBER: CPI-012CP5
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                    Sequence 7, Application US/09305923A Patent No. 6355473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Klein, Christine A.
APPLICANT: Murphy, Andrew J. M.
TITLE OF INVENTION: Methods and
TITLE OF INVENTION: Identifying
NUMBER OF SEQUENCES: 98
                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (617)227-740
TELEFAX: (617)227-4214
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 9 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , Application US/08582333A 6255059
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APPLICANT: Fowlkes, Dana M.

TITLE OF INVENTION: YEAST CELLS EXPRESSING MODIFIED G PROTEINS AND METHODS

TITLE OF INVENTION: OF USE THEREFOR

FILE REFERENCE: CPI-012CP8

CURRENT APPLICATION NUMBER: US/08/946,298A

CURRENT FILING DATE: 1997-10-07

EARLIER APPLICATION NUMBER: 08/689,172

EARLIER FILING DATE: 1996-08-06

EARLIER FILING DATE: 1996-01-17

EARLIER FILING DATE: 1996-01-17

EARLIER APPLICATION NUMBER: 08/582,333

EARLIER FILING DATE: 1995-06-05

EARLIER APPLICATION NUMBER: 08/463,181

EARLIER APPLICATION NUMBER: 08/463,181

EARLIER APPLICATION NUMBER: 08/463,181

EARLIER FILING DATE: 1995-06-05

EARLIER FILING DATE: 1995-06-05

EARLIER FILING DATE: 1994-10-13

EARLIER FILING DATE: 1994-10-13
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SEQ ID NO 4
LENGTH: 9
TYPE PRT
ORGANISM: Saccharomyces cerevisiae
US-08-946-298-4
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                                                                                                                                                                                                                                                                                                                      RESULT 15
US-09-217-609A-8
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US-08-946-298-4
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; TYPE: PRT
; ORGANIEM: Saccharomyces cerevisiae
US-09-305-923A-7
                                                                                                                                                            Patent No. 6071733

GENERAL INFORMATION:

APPLICANT: MURAMATS

APPLICANT: KOIKE, F

APPLICANT: COURA, F

APPLICANT: KOYAMA,
                                                                                                                                                                                                                                                                              Sequence 8, Application US/09217609A Patent No. 6071733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 55.6%;
Best Local Similarity 33.3%;
Matches 3; Conservative 9
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Best Local Similarity 33.3%;
Matches 3; Conservative
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EARLIER FILING DATE: 1994-09-20
EARLIER FILING DATE: 1994-01-31
EARLIER FILING DATE: 1994-01-31
EARLIER FILING DATE: 1994-01-31
EARLIER APPLICATION NUMBER: 08/041,431
EARLIER FILING DATE: 1993-03-31
NUMBER OF SEQ ID NOS: 86
                                                APPLICANT: SHIMIZÜ, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: 1025 Connecticut Avenue, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 VLHDDLLEA 9
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1 LIHEDIAKA 9
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1 LIHEDIAKA 9
                                                                                                               MURAMATSU, Masayoshi
KOIKE, Ayumi
OGURA, Kyozo
KOYAWA, Tanetoshi
SHIMIZU, Naoto
CHO, Yenwin
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
5; Mismatches 1
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Search completed: April Job time : 28.6667 secs
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PRIOR APPLICATION NATA:
APPLICATION NUMBER: 08/873,23
FILING DATE: 11-Jun-1997
AUTORNEY/AGENT INPORMATION:
NAME: TOPPENETTI, Judith L.
REGISTRATION NUMBER: 39,048
REFERENCE/DOCKET NUMBER: 1023.
TELECOMMUNICATION INPORMATION:
TELEPAN: 202-429-1776
TELEPAN: 202-429-0796
INFORMATION POR SEQ ID NO: 8:
                                                                                                                                                    Matches
                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: DC
COUNTRY: US
ZIP: 20036
ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/217,609A
FILING DATE:
CLASSITEDSATION.
                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
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 Conservative

                   6, 2006, 16:27:38
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Pred. No. 4.6e+05;
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Gaps

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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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    seq length: 0
seq length: 9
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Match Length
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Gapop 10.0 , Gapext 0.5
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US-10-623-176-2
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US-10-623-176-41
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US-10-623-176-10
US-10-623-176-10
US-10-623-176-40
US-10-623-176-14
US-10-623-176-14
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US-10-623-176-14
US-10-623-176-14
US-10-623-176-15
US-10-623-176-15
US-10-623-176-15
US-10-623-176-15
US-10-623-176-13
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2, Appli
1, Appli
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T 2 -791-: nuence licat: ERAL: PLICAL PLICAL PLICAL		Query Matcl Best Local Matches	OTHER INF FEATURE: NAME/KEY: LOCATION: 0-623-176	LENGTH: 9 TYPE: PRT ORGANISM: FEATURE:		PRIOR A	PRIOR A	FILE RECURRENT	APPLICANT: APPLICANT: TITLE OF IN	ULT 1 10-623-176- equence 2, ublication ENERAL INFO APPLICANT:		υ <u>1</u> 2 ι	- Ki F	800	2 9 Q.	100	28 29 31
USSULT 2 15-10-791-217-2 Sequence 2, Applicat Publication No. US20 GENERAL INFORMATION: APPLICANT: GOULINY, APPLICANT: Hunt, D APPLICANT: Engelha	1 1 \$=\$. 5	ı o	I: 9 PRT SM: Art	ILING DEC	ILING PPLICA	PPLICA	REFERENCE: ENT APPLICA ENT FILING	Z C			221	225	222	222	22 2 E	0 2 2 2 3 A 6 A 6 A 6 A 6 A 6 A 6 A 6 A 6 A 6 A
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ion 040 Bls	VIHDDLLEA 9	Similarity 9; Conservat		icial s	2000 2000 Ver.	N NUMBER: PCT E: 1998-07-23	R APPLICATION NUMBER: 09/489,7 R FILING DATE: 2000-01-21 R APPLICATION NUMBER: EP 97202 B FILING DATE: 1997-07-23	FERENCE: 2183-6047US APPLICATION NUMBER: FILING DATE: 2003-	Hunt, Donald F. Engelhard, Victor ENTION: HA-1 epit	cation US/1062 S20040092446A1 ON:		ພໍພໍເ	9 9 9				
US/107 91268A A.J.M d F Victor		100 100 ive	crij	eque	ن د	- R:	- R - C - R - C - C - C - C - C - C - C	47U BER 003	d F. Victor 1 epit	US/ 924		ωω.	1	φφφ	9 7 6 6	νφοι	νοοσ
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US-10-791-217-2
                                                                                                                                                                                                          RESULT 4
US-11-007-740-20
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; OTHER INFORMATION: HA-1 peptide
US-10-861-335-1
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             Sequence 20, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulimy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
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Best Local Similarity
Matches 9; Conserv
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/861,335
CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR PILING DATE: 2001-12-05
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goulmy, Elsa A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot
TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
PRIOR FILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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RESULT 6
US-10-623-176-45
; Sequence 45, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
   APPLICANT: Goulmy, Els A.J.M.
; APPLICANT: Hunt, Donald F.
; APPLICANT: Engelhard, Victor H.
; TITLE OF INVENTION: HA-1 epitopes and uses thereof; FILE REFERENCE: 2183-6047US; CURRENT APPLICATION NUMBER: US/10/623,176
; CURRENT FILING DATE: 2003-07-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Goulmy, B18 A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLS OF INVENTION: HA-1 epitopes and uses the
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP97202303.0
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-2
PRIOR PILING DATE: 1998-07-2
PRIOR FILING DATE: 1998-07-2
PRIOR FILING DATE: 2000-01-24
NUMBER: DESCRIPTION NUMBER: DE 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER: DESCRIPTION UNMBER: DE 2000-504165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 41, Application US/10623176
Publication No. US20040092446A1
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: SITE
LOCATION: (1)..(9)
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nes 8; Conserv
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100.0%; Pred. No.
cive 0; Mismatcl
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Pred. No. 1.7e+06;
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PRIOR APPLICATION NUMBER: EP 97202303.0 PRIOR FILING DATE: 1997-07-23

APPLICATION NUMBER: 09/489,760 FILING DATE: 2000-01-21

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Best Local Similarity
""" thes 8; Conserve
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-45
                                                                                                                                         ; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-10
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PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
                                                                                        Query Match
Best Local (
                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald P.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 45
LENGTH: 9
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PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
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CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                   LENGTH: 9
TYPE: PRT
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                                                                                                                                                                                               ORGANISM: Artificial Sequence FRATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE:
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                                                                   Similarity
8; Conserv
                               VIHDDILEA 9
VLRDDLLEA 9
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No. US20040092446A1
                                                                   Conservative
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                                                                                    82.2%;
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Pred. No. 1.7e+06;
0; Mismatches 1; Indels
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RESULT 8

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US-10-623-176-47
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APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 37202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER: OF SEQ ID NOS: 77
NUMBER: DESCRIPTION UNBER: PCT/NL98/004165
PRIOR FILING DATE: 2000-01-24
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                                                                                                                                                         ; OTHER INFORMATION: obtained from KIAA0223 partial complementary US-10-791-217-5
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US-10-791-217-5
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
ITILE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
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Best Local S
Matches 8
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LENGTH: 9
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SEQ ID NO 47
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                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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LOCATION: (1)...
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1 VLHDDLLEA 9
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8; Conser
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                                                                                       82.2%;
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100.0%; Pred. No. 1.7e+06;
tive 0; Mismatches 0;
                                                                   Score 37; DB 4; Le
Pred. No. 1.7e+06;
0; Mismatches 1;
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                Query Match
Best Local Similarity
"hehes 8; Conserve
                                                                                                                   US-10-623-176-1
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APPLICANT: GOULMY, E18A
APPLICANT: GOULMY, E18A
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REPERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
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Publication No. US20040092446A1
GENERAL INFORMATION:
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
FITE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT EPPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
                                                                                                                                                                                                                                                                               LENGTH: 9
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                                                                                                                                PEATURE:
NAME/KEY: SITE
LOCATION: (1)..(9)
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                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide OTHER INFORMATION: wherein X can be R or H
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1 VLHDDLLEA 9
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                                     Conservative
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Pred. No. 1.7e+06;
0; Mismatches 1
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                                                      Score 36; DB 4;
Pred. No. 1.7e+06;
                                       Mismatches
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GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/99/489,760
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR PILING DATE: 1908-07-23
NUMBER OF SEQ ID NOS: 17
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US-10-791-217-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: EP 0720303.0
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
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SEQ ID NO 40
                                                                                                                                                                                      SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10791217 Publication No. US20040191268A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                   LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: SITE LOCATION: (1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (3)...(3)
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                                                                   FEATURE:
OTHER INFORMATION: obtained from histocompatibility antigen
                                                                                                                ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 9
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100.0%; Pred. No. 1.
tive 0; Mismatches
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Sequence 42, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT PILING DATE: 2003-07-18
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 1907-07-23
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: POT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: DET/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: DET/NL98/00424
PRIOR FILING DATE: 1908-07-23
PRIOR APPLICATION NUMBER: DET/NL98/00424
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 42
LENGTH: 9
TYPE: PRT
TYPE: PRT
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; NAME/KEY: MISC FEATURE
; LOCATION: (3):...(3)
; OTHER INFORMATION: Xaa represents a histidine (H) or an arginine (R) residue
US-11-007-740-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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US-11-007-740-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
PILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR PILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIn version 3.1
SEQ ID NO 29
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Matches 8; Conservat
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TYPE: PRT
ORGANISM: Human
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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Pred. No. 1.7e+06;
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Maximum Match 100%
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Perfect score:
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       Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/US03_NEW_PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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1 US-11-129-143-163

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1 US-11-129-143-161

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1 US-11-045-788A-353

1 US-10-485-788A-355

1 US-11-033-039-785

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169, App
169, App
1785, App
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3785, App
5444, App
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5445, App
5467, Ap
5785, Ap
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Qy 1 VIHDDILEA 9 Db 1 VIHDDILEA 9 RESULT 2 US-11-010-748A-12 Sequence 12, Application US/11010748A Publication No. US20050244421A1 GENERAL INFORMATION: APPLICANT: Merck Patent GmbH APPLICANT: STRITTMATTER, Wolfgang APPLICANT: MOLL, Heidrun APPLICANT: SCHARM, Burkhard	Query Match 100.0%; Score 45; DB 7; Length 9; Best Local Similarity 100.0%; Pred. No. 1.4e+05; Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	INGTH: (PE: PF (GANISN INTURE: INTU	RESULT 1 US-11-010-748A-11 US-11-010-748A-11 US-11-010-748A-11 Sequence 11, Application US/11010748A Publication No. US20050244421A1 Publication No. US20050244421A1 APPLICANT: Merck Patent GmbH APPLICANT: MCLL, Heidrun APPLICANT: MOLL, Heidrun APPLICANT: SCHARM, Burkhard TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE PILE REFERENCE: MER-136 CURRENT FILING DATE: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE PRICE REFIGURED DATE: 2003-06-13 PRICE APPLICATION NUMBER: BCT/EF03/06251 PRICE APPLICATION NUMBER: BC02013423.5 PRICE FILING DATE: 2003-06-13 PRICE FILING DATE: 2003-06-13 PRICE FILING DATE: 2003-06-13 NUMBER OF SEQ ID NOS: 926 SOFTWARE: PatentIn version 3.1	ALIGNMENTS	26 21 46.7 4 6 US-10-485-788A-351 Sequence 28, Appl 28 20 44.4 4 6 US-10-657-295-263 Sequence 263, Appl 29 20 44.4 4 7 US-11-019-027-5 Sequence 264, Appl 30 20 44.4 4 7 US-11-1019-027-5 Sequence 264, Appl 31 20 44.4 4 7 US-11-1019-027-5 Sequence 64, Appl 32 24.4 4 7 US-11-103-030-3 Sequence 65, Appl 33 20 44.4 5 7 US-11-103-103-3 Sequence 168, Appl 36 20 44.4 5 7 US-11-129-143-168 Sequence 168, Appl 36 20 44.4 5 7 US-11-129-143-169 Sequence 169, Appl 37 20 44.4 5 7 US-11-129-143-170 Sequence 170, Appl 39 20 44.4 5 7 US-11-129-143-170 Sequence 171, Appl 39 20 44.4 5 7 US-11-129-143-171 Sequence 172, Appl 39 20 44.4 5 7 US-11-129-143-171 Sequence 172, Appl 39 20 44.4 5 7 US-11-129-143-173 Sequence 173, Appl 40 20 44.4 5 7 US-11-129-143-173 Sequence 173, Appl 41 20 44.4 6 6 US-10-857-435A-367 Sequence 367, Appl 42 20 44.4 6 6 US-10-857-435A-367 Sequence 37, Appl 43 20 44.4 6 6 US-10-857-435A-367 Sequence 67, Appl 43 20 44.4 8 7 US-11-021-305-65 Sequence 65, Appl 45 20 44.4 8 7 US-11-021-305-65 Sequence 65, Appl 5 Sequence 65, Appl 5 Sequence 65, Appl 5 Sequence 65, Appl 65 Sequence 66, A
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APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, We
                                                                                                 Sequence 164, Application US/11129143
Publication No. US20050266518A1
GENERAL INFORMATION:
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Best Local
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SEQ ID NO 163
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Best Local
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CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
PRIOR APPLICATION NUMBER: PCT/EP03/06251
PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR APPLICATION NUMBER: E902-06-13
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
          APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
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TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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COTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by TOTHER INFORMATION: ell epitopes
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ORGANISM: Homo sapiens
 HPLICANT:
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8; Conserv
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HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
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YELISEEV, Alexei A.
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Pred. No. 1.4e+05;
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SEQ ID NO 166 LENGTH: 6

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GENERAL INFORMATION:
APPLICANT: BERRY, Alan
APPLICANT: BERTZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: HUMBELIN, Markus
APPLICANT: HOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEBY, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION NUMBER: US/11/129,143
NUMBER OF SEQ ID NOS: 197
SOPTWARE: Patentin version 3.1
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US-11-129-143-165
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CURRENT FILING DATE: 2005-05-13
CURRENT FILING 
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Matches
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Publication No. US20
GENERAL INFORMATION:
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SEQ ID NO 165
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Best Local Similarity
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APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
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CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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ORGANISM: Escherichia coli
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Pred. No. 1.4e+05;
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Pred. No.
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                                                                                                                               US-11-129-143-162
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                             Query Match
Best Local Similarity 80.0
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Best Local Similarity 80.0
                                                                                                                                                                                             APPLICANT: MAYER, Anne F.
APPLICANT: VELISEEV, Alexei A.
APPLICANT: VELISEEV, Alexei A.
APPLICANT: VELISEEV, Alexei A.
APPLICANT: VELISEEV, Alexei ISOPRENOID PRODUCTION
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOPTWARE: Patentin version 3.1
SEQ ID NO 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 162, Application US/11129143
Publication No. US20050266518A1
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Best Local Similarity
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SEQ ID NO 161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: YELISEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BERRY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                             TYPE: PRT
ORGANISM: Rhizobium
                                                                                                                                                                                       LENGTH:
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2 LHDDL 6
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HUMBELIN, Markus
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MAYER, Anne F.
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                                                                                                                                                                                                                                                                                                                                                                                                                               Alan
                                                                                                                                               sp. strain NGR234
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80.0%;
                                                                       55.6%;
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                                                    Score 25; DB 7; Le
Pred. No. 1.4e+05;
1; Mismatches 0;

    Mismatches

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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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                                                                                          Length 6;
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CURRENT EILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOPTWARE: PATENTIN VETBION 3.1
SEQ ID NO 353
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                                                                                                                                                                j TYPE: PRT
j ORGANISM: Homo sapiens
US-10-485-788A-353
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US-11-004-399-1155
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Best Local S
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Best Local Similarity 100.
The A; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ramalingam, Sathiskumar
APPLICANT: Poon, Leo Lit Man
APPLICANT: Petris, Joseph Sriyal Malik
TITLE OF INVENTION: Genetically Modified Plants Comprising SARS-COV Viral Nucleotide
TITLE OF INVENTION: Sequences and Methods of Use Thereof For Immunization Against S
FILE REFERENCE: 2589/73166/RDK
CURRENT APPLICATION NUMBER: US/11/004,399
CURRENT FILING DATE: 2004-12-03
PRIOR APPLICATION NUMBER: US 60/527,637
PRIOR APPLICATION DATE: 2003-12-03
NUMBER OF SEQ ID NOS: 4043
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions
FILE REFERENCE: 20054-003320US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lu, Peter S. APPLICANT: Rabinowitz, APPLICANT: Schweizer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Chye, Mee Lee
APPLICANT: Li, Hong Ye
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                                                                                                                                                                                                                      ENGTH: 6
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                 1 LVHDDV 6
                                                      1 VLHDDL 6
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3; Conserv
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                                                                                                        51.1%;
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                                                                                       Score 23; DB 6;
Pred. No. 1.4e+05;
3; Mismatches 0
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Gaps

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Gaps

US-10-485-788A-354

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PRIOR APPLICATION UMBER: WO PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: Patentin version 3.1
SEQ ID NO 355
LENGTH: 8
TYPE: PRT
ORGANISM: Homo sapiens
US-10-485-788A-355
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; ORGANISM: Homo sapiens
US-10-485-788A-354
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Best Local Similarity 50.0
Matches 3; Conservative
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                                              Matches
                                                               Best Local Similarity
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CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 20054-003320US
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2002-00 NUMBER OF SEQ ID NOS: 841
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APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
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1 VLHDDL 6
                                              Conservative
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Carrick, Deanna Marie
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                                          51.1%; Score 23; DB 6; Le 50.0%; Pred. No. 1.4e+05; tive 3; Mismatches 0;
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Pred. No. 1.4e+05;
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Sequence 809, Application US/11033039

publication No. US20060002947A1

GENERAL INFORMATION:
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
TITLE OF INVENTION: LI-KRY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01

CURRENT APPLICATION NUMBER: US/11/033,039

CURRENT FILING DATE: 2005-01-11

PRIOR APPLICATION NUMBER: 10/245,871

PRIOR APPLICATION NUMBER: 10/245,871

PRIOR APPLICATION NUMBER: 10/197,000

PRIOR FILING DATE: 2002-09-17

PRIOR FILING DATE: 2002-09-17

PRIOR APPLICATION NUMBER: 09/396,813

PRIOR FILING DATE: 2002-09-14

NUMBER OF SEQ ID NOS: 1452

SOFTWARE: PatentIn version 3.3

SEQ ID NO 809
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US-11-033-039-785
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Publication No. US20060002947A1
GENERAL INFORMATION:
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.3 SEQ ID NO 785
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Best Local Similarity 44.4%;
Matches 4; Conservative
                                                                                          Matches
                                                                                                          Query Match
Best Local Similarity 44.4%;
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TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 09/396,813
PRIOR FILING DATE: 1999-09-14
NUMBER OF SEO ID NOS: 1452
COLUMN OF SEO ID NOS: 1452
                                                                                                                                                                                                                                 LENGTH: 9
TYPE: PRT
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LLHESTLKA 9
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Pred. No. 1.4e+05;
3; Mismatches 2
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Pred. No. 1.4e+05;
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RESULT 15 US-10-485-788A-352 ; Sequence 352, Application US/10485788A

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GENERAL INFORMATION:
APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
APPLICANT: Schweizer, Johannes
APPLICANT: Schweizer, Johannes
APPLICANT: ARbor Vita Corporation
FITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-0033200S
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2001-08-03
PRIOR PRILING DATE: 2002-08-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: PatentIn version 3.1
SOFTWARE: PatentIn version 3.1
SOFTWARE: PatentIn version 3.1
CNGANISM: Homo sapiens
Search completed: April 6, 2006, 16:51:13 Job time: 13.3333 secs
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1 VHDDV 5
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB
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Perfect score:
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2: geneseqp2000
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4: geneseqp2001
5: geneseqp2003
6: geneseqp2003
7: geneseqp2003
8: geneseqp2003
9: geneseqp2003
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geneseqp20008:*
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geneseqp20048:*
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Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	S	4	ω	N	,_	Result No.
23	23	23	23	23	23	24	24	24	25	25	25	26	26	27	29	36	36	37	37	37	42	42	42	Score
54.8	54.8	54.8	54.8	54.8	54.8	57.1	57.1	57.1	9	59.5	59.5	61.9		64.3	69.0	85.7	85.7	88.1	88.1	88.1	100.0	100.0	100.0	Query Match I
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ABR19174	ABR19578	ABR18988	ABR19426	ABR19981	AAU71211	ADM18218	ADM18256	AAB81550	ADT73689	ADT72766	AAU71428	ADT72767	ADT73690	AAW47427	ADT73023	AAW97572	AAW99195	ADH40333	AAW97375	AAW99196	ADH40334	AAW97374	AAW99197	ID
Abr19174 Human can	Abr19578 Human can	Abr18988 Human can	Human	Abr19981 Human can	Aau71211 Human MHC	Adm18218 Chlamydia	Adm18256 C trachom	Aab81550 Zif268 zi			Aau71428 Human MHC	Adt72767 Human RSV	Adt73690 Human RSV	Aaw47427 Prenyl di	Adt73023 Human RSV	Aaw97572 T-cell ep	Aaw99195 Minor his	Adh40333 Human min	Aaw97375 HA-1 H-al	Aaw99196 Minor his	Adh40334 Human min	Aaw97374 HA-1 R-al	97 Minor h	Description

4 5	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	
21	21	21	21	21	21	21	21	21	21	21	21	21	22	22	22	22	23	23	23	
50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	52.4	52.4	52.4	52.4	54.8	54.8	54.8	
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ADM20449	ADA62245	ADA62237	ADA62240	ABP96179	ABG79371	ABP48419	ABP48414	ABP48411	ADE84629	AAU80810	AAB12035	AEB31418	ADW23047	ABY01413	ADF72091	AAW97373	ADZ50925	ABY01546	ADU99853	***************************************
Adm20449	Ada62245	Ada62237	Ada62240	Abp96179	Abg79371	Abp48419	Abp48414	Abp48411	Ade84629	Aau80810	Aab12035	Aeb31418	Adw23047	Aby01413	Adf72091	Aaw97373	Adz50925	Aby01546	Adu99853	
Synthetic	Zinc fing		Zinc fing	Zinc fing	CXCR zinc	Zinc fing	Zinc fing	Zinc fing	Mammalia	Rat Rb-in	Peptide #	Endoplasm	SARS coro	SARS coro	Rabbit pe	Peptide e	Y. pestis	SARS coro	BCZ4 tumo	

ALIGNMENTS

RESULT 1
AAW99197
ID AAW9
XX AAW9
AC AAW9
XX Mino
XX WPI
PH 04-F
PF 23-7
XX WPI
XX Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease. Minor histocompatibility antigen HA-1 T-cell epitope #3 23-JUL-1997; 23-JUL-1998; 04-FEB-1999. WO9905174-A1. Homo sapiens. 20-MAY-1999 AAW99197; AAW99197 standard; peptide; 9 AA. (UYLE-) RIJKSUNIV LEIDEN. (first entry) 97EP-00202303. 98WO-NL000425.

A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.

WPI; 1999-153312/13. Goulmy EAJM, Hunt DF,

Engelhard VH;

Disclosure; Page 15; 47pp; English.

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the diseases treatment of severe aplastic anaemia, leukaemia, and immune deficiency

Sequence 9 AA;

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ARM97137 2
AAM973 AAM973 AAM973 AAM973 AAM973 AAM99 AXX
AAM97 AAM99 AXX
AAM97 AAM99 AXX
AAM97 AAM99 AXX
AAM97 AAM99 AAM97 AAM999 AAM99 AAM999 AAM999 AAM999 AAM999 AAM999 AAM999 AAM999 AAM999 AAM999 
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RESULT 3
ADH40334
ID **ADH40334 standard; peptide; 9 AA.
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AC ADH46034;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
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02-JUN-1998;
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9; Conserv
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98EP-00870125.
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Pred. No. 2e+06;
Mismatches 0
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Pred. No.
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Matches
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Minor histocompatibility antigen; HA-1; T-cell epitope; immuno graft versus host disease; bone marrow transplant; leukaemia; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
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                                                                  histocompatibility antigen HA-1 T-cell epitope #2.
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9; Conserv
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Pred. No. 2e+06;
Mismatches
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RESULT 5
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WPI; 1999-142960/12
                                                               23-JUL-1997;
02-JUN-1998;
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                                                                                                                                                              Homo
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                                                                                                                                                                                                                             HA-1 H-allele sequence.
                                                                                                                                                                                                                                                                                            AAW97375 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
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                                                                                              23-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9 AA;
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                                         RIJKSUNIV LEIDEN
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                                                              97EP-00202303
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                                                                                                                                                                                                                                                                                            protein; 9 AA.
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2e+06;
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RESULT 6
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The invention relates to a novel method for providing epitopes of allelic variants of antigenic proteins from specific species based on single nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein, identifying, selecting allelic peptide/protein variants, expression product or its fragment encoded by DNA sequence having SNP, creating variant epitopes, selecting epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is
                                                                                                                                                                                                       Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                              WPI; 2004-082200/08
                                                                                                                                                                                                                                                                                                              Strittmatter W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Typing minor histocompatibility antigen HA-1 - by amplifying identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. of genetic aberrances.
                                                                                                                                                         Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                  13-JUN-2003; 2003WO-EP006251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003106692-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human minor histocomptibility antigen HA-1 T cell eptiope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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8; Conserv
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88.9%;
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Pred. No.
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2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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Synthetic.
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                                                          The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                diseases
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Sequence 9 AA;
                                                                                                                                                                                                                    A new minor histocompatibility antigen, diseases and prevent rejection and host
                                                                                                                                                                                                                                                                                                  Goulmy EAJM,
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                                                                                                                                                                          Claim 1; Page 32; 47pp;
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                                                treatment of severe aplastic
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8; Conserv
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Pred. No. 2e+C
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                                                anaemia,
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2e+06;
                                                   leukaemia,
                                                                                                                                                                                                                       HA-1 - useful to treat immune versus graft disease in bone
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                                                   and immune
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                                                                                                                                                                                                                                                                                                          The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides.
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                                                                                                                                                                                                          Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 39; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 versus-host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9905173-A1
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                                                    1 VLRDDLLEA 9
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8; Conserv
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Pred. No. 2e+06;
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RESULT 9 ADT73023 ID ADT7

ADT73023 standard; peptide; 9 AA.

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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                  This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins non-structural proteins NS-1 (also CC known as the IC protein) & NS-2 (1B protein), a polymerase protein I and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC composition comprising an above peptide mixed with a pharmaceutical cc excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC RSV peptide is useful for preparing a diagnostic composition or an RSV (prophylactic or therapeutic) vaccine composition for a DNA-based CC (mmunisation, or for preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the polypeptide in an immune response-provoking amount of specific CT1).

CC accordingly, these peptide compositions have virucidal activity. This computation is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                         Query Match
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Matches 4
 Prenyl diphosphate synthetase preserved region VI sequence.
                                   05-JUN-1998
                                                                     AAW47427;
                                                                                                                                                                                                                                                                                                                            Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                 AAW47427 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 16; SEQ ID NO 919; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human respiratory syncytial virus; cytotoxic T lymphocyte; CTL; MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence from preserved region VI of prenyl diphosphate synthetase (PDS), was used in the preparation of primers for the amplification of PDS DNA. Substances synthesised by PDS, i.e. polyprenyl diphosphates, are precursors of physiologically active substances, e.g. vitamin K and ubiquinones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA encoding prenyl diphosphate synthetase subunit(s) - new Micrococco prenyl diphosphate synthetase subunit polypeptide(s), and methods for preparing enzymes from subunit(s).
                  WPI; 2004-758334/74.
                                                                                                                                                              Human respiratory syncytial virus
                                                                                                                                                                                   human respiratory syncytial virus; cytotoxic T lymphocyte; MHC epitope; DNA-based immunisation; RSV infection: virucic
                                                                                                                                                                                                                 Human RSV L high affinity binding peptide SeqID 1586
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9
                                                                                                   16-APR-2004; 2004WO-EP004061.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JUN-1996;
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                                       Lasters
                                                                              16-APR-2003; 2003EP-00447095
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New respiratory syncytial virus (RSV) peptides (e.g.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC eight structural proteins protein). Specifically, it refers to a CC composition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a mucleic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC Tlymphocytes (CTLs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (prophylactic or therapeutic) vaccine composition for a DNA-based CC (prophylactic or therapeutic) vaccine being prepared by contacting the conjugatide in an immune response-provoking amount of specific CTL). CC coordingly, these peptide compositions have virucidal activity. This coppide sequence is a human RSV high binding affinity peptide of the
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                                                         Disclosure; SEQ ID NO 663; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADT72767 standard; peptide; 9
                                                                                                          New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
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based immunisation; RSV infection; virucidal; vaccine
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al virus (RSV), in particular
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This invention relates to novel isolated or purified peptides of the

Sequences AAU71093-AAU71796 represent the 103P3E8-related protein epptide fragments of the protein. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expression.

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Disclosure; Page 90; 128pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human respiratory syncytial virus (RSV), in particular ten RSV genes encoding 11 separate viral proteins: non-structural proteins NS-1 (a. known as the 1C protein) & NS-2 (1B protein), a polymerase protein Leight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, SH (also known as the 1A protein). Specifically, it refers to a composition comprising an above peptide mixed with a pharmaceutical excipient or an RSV immunogenic composition comprising a recombinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   103P3E8; prostate; bladder; kidney; colon; lung; breast; rectum; stomach; tumour; cancer; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human MHC
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                                                                                                             Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene
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RESULT 14
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This invention relates to novel isolated or purified peptides of the human respiratory syncytial virus (RSV), in particular ten RSV genes encoding 11 separate viral proteins: ROV), in particular ten RSV genes encoding 11 separate viral proteins (RSV), in particular ten RSV genes encoding 11 separate viral proteins: RSV), in particular ten RSV genes encoding 11 separate viral proteins (RSV), in particular protein RS-1 (also known as the 1A protein). Specifically, it refers to a composition comprising an above peptide mixed with a pharmaceutical excipient or an RSV immunogenic composition comprising a recombinant excipient or an RSV immunogenic composition comprising a recombinant excipient or with a nucleic acid insert encoding an above peptide. The present invention describes an in vitro method of detecting cytotoxic 1 lymphocytes (CTLs) that respond to a major histocompatibility complex (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The RSV peptide is useful for preparing a diagnostic composition or an RSV (prophylactic or therapeutic) vaccine composition for a DNA-based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 1019388 polymucleotide, its relate protein and peptide fragments and specific PCR primers are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 1039388-related protein, and a ribozyme capable of cleaving a polymucleotide having the 1039388 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 1039388. The sequences can be used in diagnostic methods to monitor the level of 1039388 gene products in serum, blood, urine and tissue and to thereby detect the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
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ADT73689
ID ADT7
   CC This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC SH (also Known as the 1A proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC composition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC Thymphocytes (CTLs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (prophylactic or therapeutic) vaccine composition for a DNA-based CC (prophylactic or therapeutic) vaccine being prepared by contacting the CC immunisation, or for preparing an immune response provoking vaccine in Cthe event of RSV infection (the vaccine being prepared by contacting the polypeptide in an immune response-provoking amount of specific CTL).

CC polypeptide in an immune response provoking vaccine in composition in a immune response provoking and activity. This composition is a human RSV high binding affinity peptide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                      Claim 16; SEQ ID NO 1585; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-758334/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lasters I, Desmet J,
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		Search completed: April 6, 2006, 16:19:10 Job time : 112.667 secs	Oy 1 VLRDDLL 7 ::: : bb 2 IIKDDLL 8	Ouery Match 59.5%; Score 25; DB 8; Length 9; Best Local Similarity 42.9%; Pred. No. 2e+06; Matches 3; Conservative 4; Mismatches 0; Indels 0; Gaps 0;	
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GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: April 6, 2006, 16:19:36; Search time 18.6667 Seconds (without alignments) 46.390 Million cell updates/sec

Title: Perfect score: Sequence: US-10-791-217A-5 42 1 VLRDDLLEA 9

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters:

791

Minimum DB seq length: 0
Maximum DB seq length: 9

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : PIR 80:*
1: pir1:*
2: pir2:*
3: pir3:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ribosomal protein	glycoprotein compo	hypothetical prote	neuropeptide Antho	protein-glutamine	gamma subunit of P	calliFMRFamide 10	neuropeptide Grb-A	neuropeptide Grb-A	transaldolase (EC	glycine cleavage s	dextransucrase (EC	chlorophyll a/b-bi	conopressin S - co	tetrameric protein	ferredoxin a2 - Ja

ALIGNMENTS

Query Match 38.1%; Score 16; DB 2; Length 8; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;		C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 31-Dec-2004 C;Accession: PC4131 R;Kawasaki, S.; Arai, H.; Igarashi, Y.; Kodama, T. Gene 167, 87-91, 1995 A;Title: Sequencing and characterization of the downstream region of the genes encoding v for blosynthesis of heme d1.	RESULT 2 PC4131 hypothetical protein 8 [imported] - Pseudomonas aeruginosa (fragment) C;Species: Pseudomonas aeruginosa	Qy 4 DDL 6 	Query Match 38.1%; Score 16; DB 2; Length 7; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	A;ACCEBBION: SERUY A;Status; preliminary A;Molecule type: protein A;Residues: 1-7 <edg> A;Cross-references: UNIPARC:UPI000017C164</edg>	R;Edgar, P.F. FEBS Lett. 375, 159-161, 1995 A;Title: Hucolin, a new corticosteroid-binding protein from human plasma with structural A;Reference number: S68004; MUID:96087107; PMID:7498469	RESULT 1 \$68004 hucolin, 75K chain - human (fragment) C;Species: Homo sapiens (man) C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999 C;Accession: \$68004	
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elastase - Pseudomonas aeruginosa (;Species: Pseudomonas aeruginosa C;Date: 19-Mar-1997 #sequence_revision 24-Jul-1997 #text_change 24-Jul-1997 C;Accession: $20446 R;Kessler, E.; Safrin, M.; Peretz, M.; Burstein, Y. FEBS Lett. 299, 291-293, 1992 A;Reference number: $20446; MUID:92183956; PMID:1544509 A;Accession: $20446
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S20446
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submitted to the Protein Sequence Database, J
A:Description: Purification of PKCI from rat
                                                                                                                                                                                     macrophage cytotoxicity-inducing factor, 29K - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
C;Accession: A60427
R;Jones, C.M.; Prince, C.A.; Williams, J.S.
Exp. Hematol. 19, 704-709, 1991
A;Title: Purification and amino acid analysis of a human macrophage cytotoxicity-inducin A;Reference number: A60427; MUID:91372335; PMID:1909970
A;Accession: A60427
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A; Accession: A59489
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C;Species: Rattus norvegicus
C;Date: 25-Aug-2003 #sequence_revision 25-Aug-2003 #text_change 25-Aug-2003
C;Accession: A59489
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                                                                                             A;Cross-references: UNIPROT:Q7M4R5; UNIPARC:UPI0000142BE0 A;Note: the sequence from the text on page 706 is inconsistent C;Keywords: cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: protein A; Residues: 1-7 < KES> A; Cross-references: UNID:
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Pred. No.
                  0;
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Pred. No. 2.8e+05;
2; Mismatches 0
                                Score 14; DB 2;
Pred. No. 2.8e+05;
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R;Gold, D.P.; Offner, H.; Sun, D.; Wiley,
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        biotin A - Citrobacter freundii (fragment)
C;Species: Citrobacter freundii
C;Species: Citrobacter freundii
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C;Accession: I40697
R;Shiuan, D.; Campbell, A.
Gene 67, 203-211, 1988
A;Reference number: I40697; MUID:89006280; PMID:2971595
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-9 <GOL>
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C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0679; PT0708
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                                                                                                                 J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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A; Residues: 1-4 < RES>
                                      A; Molecule type: DNA
A; Residues: 1-5 < FEE>
                                                                                                                                                                       R; Feeney, A.J
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A;Cross-references: UNIPARC:UPI000017C84D
A;Experimental source: day 18 fetal thymus,
                                                                            A; Status: translation not
                                                                                               A; Accession: PT0679
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100.0%; Pred. No.
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  strain BALB/c, 154-2J
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2.8e+05;
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Cerebellar degeneration-related protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 28-Sep-1990 #sequence_revision 28-Sep-1990 #text_change 24-Jun-1993
C;Accession: B35640
R;Chen, Y.T.; Rettig, W.J.; Yenamandra, A.K.; Kozak, C.A.; Chaganti, R.S.K.; Posner, J.F.
Proc. Natl. Acad. Sci. U.S.A. 87, 3077-3081, 1990
A;Title: Cerebellar degeneration-related antigen: a highly conserved neuroectodermal manaly Reference number: A35640; MUID:90222173; PMID:2326268
A;Accession: B35640
                                                                                                                                                                                                                                                                     RESULT 10
B35640
                     A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-6 <CHE>
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A;Title: Junctional sequences of fetal T cell receptor beta
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0601
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A;Experimental source: newborn thymus, strain BALB/c, 161-28
C;Keywords: T-cell receptor
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A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-5 <PE2>
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A;Experimental source: day 18 fetal thymus,
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A;Cross-references: UNIPARC:UPI000017C63A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Keywords: T-cell receptor
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A;Experimental source: newborn thymus, strain BALB/c, clo
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A; Residues: 1-5 < FEE>
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Pred. No. 2.8e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 5;
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Query Match

Score 12;

DB

2

Length 6;

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R;Cala, S.B.; Jones, L.R.
J, Biol. Chem. 266, 391-398, 1991
A;Title: Phosphorylation of cardic
A;Reference number: A39040; MUID:
A;Accession: B39040
A;Status: preliminary
                                                                                                                             RESULT 13
B39040
Calsequestrin, fast skeletal muscle - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 23-Aug-1991 #sequence_revision 23-Aug-1991 #tex
C;Accession: B39040
C;Accession: B39040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Gibney, G.; MacPhee-Quigley, K.; Thompson, B.; Vedvick, T.; Low, M.G.; Taylor, S.S.; J. Biol. Chem. 263, 1140-1145, 1988
A;Title: Divergence in primary structure between the molecular forms of acetylcholinest, A;Reference number: A34026; MUID:88087239; PMID:3335534
A;Accession: A34026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acetylcholinesterase (BC 3.1.1.7) 5.6S form - Pacific electric ray (fragment) C;Species: Torpedo californica (Pacific electric ray) C;Date: 16-Mar-1990 #sequence_revision 16-Mar-1990 #text_change 08-Nov-1996 C;Date: 16-Mar-1990 #sequence_revision 16-Mar-1990 #text_change 08-Nov-1996
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C;Keywords: alternative splicing; carboxylic ester hydrolase
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J. Exp. Med. 174, 115-124, 1991
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: protein A; Residues: 1-7 < GIB>
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A;Experimental source: adult thymus, strain BALB/c
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A; Residues: 1-6 < FEE>
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Pred. No.
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Pred. No. 2.8e+05;
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1; Mismatches 0
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2.8e+05;
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R; Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A; Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A; Reference number: PT0509; MUID:91277601; PMID:1711558
A; Accession: PT0628
A; Status: translation not shown
A; Molecule type: mRNA
A; Residues: 1-7 <FEE>
A; Cross-references: UNIPARC:UPI000017C7CD
A; Experimental source: newborn thymus, strain BALB/c
C; Keywords: T-cell receptor
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PT0628
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J. Exp. Med. 174, 115-124, 1991

A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.

A;Reference number: PT0509; MUID:91277601; PMID:1711558

A;Accession: PT0722

A;Status: translation not shown

A;Molecule type: DNA

A;Residues: 1-7 <FEE>

A;Residues: 1-7 <FEE>
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PT0722
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0628
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A;Cross-references: UNIPARC:UPI000017C5BC
C;Keywords: phosphoprotein; skeletal muscle
                                                                                                                                                                                                                                                               A;Cross-references: UNIPARC:UPI000017C81F
A;Experimental source: newborn thymus, strain BALB/c C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T-cell receptor beta chain V-D-J region (135-1G) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0722
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             Search completed: April 6, 2006, 16:26:10 Job time : 18.6667 secs
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Best Local Similarity
Matches 2; Conserv
time : 18.6667 secs
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ilarity 100.0%; Pred. No. 2.8e+05;
Conservative 0; Mismatches 0;
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Maximum Match 100%
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Perfect score:
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Maximum DB
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 9
        protein search, using sw model
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Match
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Q7KYP6_HUMAN
Q8LPT5_MAIZE
P82568_STRPY
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Q4QWYJ 9MARC
Q51594 9ZZZZ
Q8KZYA 9CHRO
Q9ZIE9 NEIME
Q7MARS HUMAN
Q9FSZZ CICAR
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Q6Q7GO RANRI
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NUCLEOTIDE SEQUENCE.
STRAIN=Y-13, and W-16;
MEDLINE=22779048; PubMed=12897253; DOI=10.1105/tpc.012526;
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9 AA. e update) ion update) ionyophyta; Tracheophyta; ; Poales; Poaceae; ; Zea.	DB 2; Length 8; 2.2e+06; 1es 0; Indels 0; Gaps 0;	10.1105/tpc.012526; Rafalski A.; quence diversity and linkage loci."; DNA.	8 AA. e update) ion update) ionyophyta; Tracheophyta; ; Poales; Poaceae; ; Zea.	Alburnum	Q6pud7 simian viru Q6pud9 simian viru Q6pue1 simian viru Q6qsx9 simian viru Q64sx4 fundulus he P82992 rhodopseudo Q7ry89 neurospora Q61cv2 homo sapien Q6uvk2 malus domes Q56se1 viburnum ur Q56se1 viburnum ur

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RESULT 3
Q705M2 HUMAN PRELIMINARY;
AC Q705M2;
DT 05-UUL-2004 (TrEMBLrel. 27,
DE Hypermethylated in cancer 1
GN Name-HIC1;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordat
OC Mammalia; Eutheria; Euarcho
OC Homo.
OX NCBI_TaxID=9606;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RC TISSUE=Mammary gland;
RA PINTESUE=Mammary gland;
RA PINTESUE=Mammary gland;
CTISSUE=Mammary gland;
RA PINTESUE=Mammary gland;
RA PINTE
RESULT 4

QTKYP6 HUMAN
ID QTKYP6;
AC QTKYP6;
DT 05-JUL-2
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OS Homo sap
OC Homo.
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RP NUCLEOTI
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pinte S., Guerardel C., Deltour S., Godwin A.K.,
"Identification of a second G-C-rich promoter cor
murine and rat tumor suppressor genes HIC1.";
Oncogene 23:4023-4031(2004).
EMBL; AJ550616; CAD79467.1; -; mRNA.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Mi.V glycoprotein (exon P3B) (Fragment).
                                                                    TISSUE=Miltenberger class MEDLINE=90005483; PubMed=
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                               Vignal A.,
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Rahuel C., El-Maliki B
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RESULT 5
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ID QBLPTS;
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AC QBLPTS;
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DT 01-OCT-2002 (TrEMBLrel. 22,
DE EMARYOCLA; VIRIGIDAL ENCOPENT
OC ENCADORATION (Maize); Magnoliophyt
OC PACCAD clade; Panicoideae; OCT NCBL TAXID=4577;
RN NUCLEOTIDE SEQUENCE.
RA Ching A.S., Caldwell K.S.,
RA MOTGENTE M., RAFALSKI J.A.;
RA MOTGENTE M., RAFALSKI J.A.;
RA EMBL; AY094310; AAM21836.1;
FT NON TER 1
SQUENCE 9 AA; 977 MW;
RESULT
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AC P8
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01-OCT-2000
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P82568 STRPY
P82568;
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Eur. J. Biochem. 184:337-344(1989)
[2]
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Blanchard C.D.;
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Vignal A., Rahuel C., El-Maliki
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NON TER

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Ching A.S., Caldwell K.S., Jung M.,
Morgante M., Rafalski J.A.;
Submitted (APR-2002) to the EMBL/Ger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota; Viridiplantae; Streptophyta; En
Spermatophyta; Magnoliophyta; Liliopsida;
PACCAD clade; Panicoideae; Andropogoneae;
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TISSUE=Miltenberger class V;
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Eur. J. Biochem. 184:0-0(1989)
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                                                                                                                Streptococcus pyogenes.
                                         NCBI_TaxID=1314;
                                                                                            Bacteria; Firmicutes;
                                                                                                                                           Unknown protein from 2D-page
PROTEIN SEQUENCE,
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  AND MASS SPECTROMETRY
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(Fragment).
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Pred. No. 2.2e+06;
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                                                                                               Streptococcaceae;
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Q80H91_9PARA
ID Q80H91_9PARA PRELIMINARY;
AC Q80H91;
DT 01-UN-2003 (TrEMBLrel. 24,
DT 01-UN-2003 (TrEMBLrel. 24,
DT 01-JUN-2003 (TrEMBLrel. 24,
DT 01-JUN-2003 (TrEMBLrel. 24,
DE HN/HNO (Fragment).
GN Name=HN/HNO;
OS Newcastle disease virus.
OC Viruses; serNA negative-str
OC Paramyxoviridae; Paramyxovi
OX NCBI_TaxID=11176;
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  Viruses; ssRNA negative-strand viruses; Paramyxoviridae; Paramyxovirinae; Avulav NCBI_TaxID=11176;
                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                      Asturias J.;
Submitted (JUL-1994) to the EMBL/GenBank/DDBJ databases.
EMBL; X80041; CAA56350.1; -; Genomic DNA.
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MEDLINE=95255652; PubMed=7737502; DOI=10.1016/0378-1119(94)00530-6;

Asturias J.A., Diaz E., Timmis K.N.;

"Evolutionary relationship of the biphenyl dioxygenase of the gram-
positive bacterium Rhodococcus globerulus P6 to multicomponent
dioxygenases of gram-negative bacteria.";

Gene 156:11-18(1995).
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Submitted (MAY-2000) to
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Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
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VanBogelen R.A.;
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MW=22592.04;
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02, Last sequence up
24, Last annotation
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Pred. No. 2.2e
4; Mismatches
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Pred. No.
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              Avulavirus
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                       Mononegavirales;
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RESULT 10
Q9XJN0_9VIRU
ID Q9XJN0;
AC Q9XJN0;
DT 01-NOV-1999 (T
DT 01-DEC-2001 (T
DT 01-DEC-2001 (T
DB P10 (Fragment)
OS Bacteriophage
OC Viruses; daRNA
OX NCBI_TaxID=908
RN [1]
RP NUCLEOTIDE SEC
RX MEDLINE=993504
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Matches
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                                             Bacteriophage phi-10.
Viruses; dsRNA viruses;
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EMBL; Y10620; CAA71631.1; -; Genomic DNA.
GO; GO:0016787; F:hydrolase activity; NAS.
SEQUENCE 9 AA; 1258 MW; C52D07340AB41412 (
                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.

MEDLINE=97336048; PubMed=9192843; DOI=10.1006/geno.1997.4731;

Riessewijk A.M., Hu L., Schulz U., Tariveridan G., Hoeglund P.,

Kere J., Ropers H.H., Kalscheuer V.M.;

"Monoallelic expression of human PEGI/MEST is paralleled by passpecific methylation in fetuses.";
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2003 (TrEMBLrel. 24,
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EMBL; 869419; AAP19628.1; -; Genomic_DNA.
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NUCLECTIDE SEQUENCE.
MEDLING=94175786; PubMed=8129624;
Collins M.S., Strong I., Alexander
"Evaluation of the molecular basis
"Evaluation of the molecular basis
  NUCLEOTIDE SEQUENCE
MEDLINE=99350412; Po
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Q9UE26;
                               NCBI_TaxID=90889;
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Mammalia; Eutheria;
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"Isolation of additional double-stranded RNA.";
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                                             Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Marchantiophyta;
Jungermanniopsida; Metzgeriidae; Metzgeriales; Pallaviciniineae;
Pallaviciniaceae; Jensenia.
NCBI_TaxID=280534;
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Forrest L.L., Schuette S.W., Crandall-Stotler "A Molecular Study of the Simple Thalloid Live (Marchantiophyta, Pallaviciniaceae).";
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Forrest L.L., Schuette
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                              NUCLEOTIDE SEQUENCE.
                                                                                                Jensenia spinosa.
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BL; AF125675; AAD22555.1; -; Genomic_RNA.
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Unknown protein from 2D-page (Fragment).
Lactobacillus sanfranciscensis (Lactobacillus sanfrancisco).
Bacteria; Firmicutes; Lactobacillales; Lactobacillus.
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EMBL; AY734726; AAX46211.1; -;
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SEQUENCE 9 AA; 1030 MW; 6D4
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                                                      MEDLINE=86223772; PubMed=2423502; Weber P.C., Palchaudhuri S.; "Incompatibility repressor in a R plasmid ColV2-K94."; J. Bacteriol. 166:1106-1112(1986)
                                                                                                                                                                                                                                                                                                                    CopB protein (Frag
Plasmid ColV2-K94.
Plasmid ColV2-K94.
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NCBI_TaxID=2458;
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Drews O., Weiss W., Reil G., Parlar H., Wait R., Goerg A.;
"High pressure effects steep-wise altered protein expression in
Lactobacillus sanfranciscensis.";
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STRAIN=DSM 20451;
EMBL; M13472;
Plasmid.
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RESULT 15

QREPX4 9CHRO

ID QREPX4, 9CHRO PRELIMINARY; PRT; 8 AA.

AC QREPX4, 1

DT 01-OCT-2002 (TrEMBLrel. 22, Created)

DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)

DT 01-OCT-2002 (TrEMBLrel. 22, Last annotation update)

DT 01-OCT-2002 (TrEMBLrel. 26, Last sequence update)

DT 01-OCT-2002 (TrEMBLrel. 27, Last sequence update)

DR McGCCYstis sp. T96-1.

GR Mame=pcA;

Microcystis sp. T96-1.

OS Microcystis sp. T96-1.

ON NCBI_TAXID-198099;

RN MCBLINE-22338268; PubMed=12450830;

RN MEDLINE-22338268; PubMed=12450830;

RN MEDLINE-2338268; PubMed=12450830;
Search completed: April
Job time : 120 secs
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Best Local Similarity 37.5%;
Matches 3; Conservative
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SEQUENCE
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NUCLEOTIDE SEQUENCE.

MEDLINE=22338268; PubMed=12450830;

DOI=10.1128/AEM.68.12.6070-6076.2002;

Baker J.A., Entsch B., Weilan B.A., McKay D.B.;

"Monitoring changing toxigenicity of a cyanobacterial bloom molecular methods.";

Appl. Environ. Microbiol. 68:6070-6076(2002).

EMBL; AY117046; AAM54719.1; -; Genomic_DNA.

NON_TER 8

8

SEQUENCE 8 AA; 890 MW; F4DB01A73771A336 CRC64;
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8 AA; 998 MW; 0F37205AA73416D7 CRC64;
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Pred. No. 2.2e+06;
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1: /cgm2_6/ptodata/1/iaa/5_COMB.pep:*

2: /cgm2_6/ptodata/1/iaa/6_COMB.pep:*

3: /cgm2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgm2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgm2_6/ptodata/1/iaa/RE_COMB.pep:*

6: /cgm2_6/ptodata/1/iaa/RE_COMB.pep:*
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US-09-217-609A-11
US-09-269-250E-38
US-08-873-23-38-11
US-09-489-760-4
US-09-586-419-14
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US-08-516-859A-20
US-08-516-859A-20
US-08-516-859A-75
US-08-516-859A-75
US-08-516-859A-75
US-08-516-472-75
US-08-516-472-75
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US-08-516-472-75
US-08-516-472-75
US-08-516-472-75
US-08-516-472-75
US-08-459-586-72
US-08-459-588-73
US-08-459-588-73
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Biocceleration Ltd.
Sequence 18, Appl Sequence 20, Appli Sequence 20, Appli Sequence 29, Appli Sequence 11, Appli Sequence 11, Appli Sequence 11, Appli Sequence 14, Appli Sequence 14, Appli Sequence 20, Appli Sequence 70, Appli Sequence 71, Appli Sequence 75, Appli Sequence 76, Appli Sequence 77, Appli Sequence 78, A
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                                                                                                                 GENERAL INFORMATION:

APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
ITITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION UNUBER: PCT/NL98/00424
EARLIER APPLICATION UNUBER: DCT/NL98/00424
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US-09-269-250B-18
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            TYPE: PRT ORGANISM:
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ALIGNMENTS	US-08-399-411-78 US-08-516-859A-72 US-08-516-859A-78 US-09-586-472-78 US-09-586-472-78 US-09-586-72 US-09-528-706-72 US-09-528-706-72 US-09-528-706-72 US-09-35-430-515 US-09-35-430-565 US-10-394-980-421 US-10-394-980-421 US-10-394-980-421 US-10-394-980-451 US-10-394-980-451 US-10-394-980-451 US-09-803-951-11
	Sequence 78, Appl Sequence 72, Appl Sequence 78, Appl Sequence 78, Appl Sequence 78, Appl Sequence 78, Appl Sequence 79, Appl Sequence 15, Appl Sequence 15, Appl Sequence 421, Appl Sequence 464, Appl Sequence 11, Appl Sequence 565, Appl Sequence 57, Appl Sequence 57, Appl Sequence 51, Appl

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; OTHER INFORMATION: Description of Artificial Sequence: Exon US-09-269-250E-18
                                  US-09-489-760-5
                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION: Blsa
APPLICANT: GOULMY, ESSA
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 18
Sequence 5, Application US/09489760 Patent No. 6878375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18, Application Patent No. 6830883
                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Pred. No. 4.6e+05;
0; Mismatches 0;
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KIAA0223 partial complementary DNA

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Sequence 2, Application US/09489760
Patent No. 6878375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
APPLICANT HARD, Victor H
TITLE OF INVENTION: The HA-1 Antigen
RESULT 5
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Patent No. 6830883

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994

CURRENT APPLICATION NUMBER: US/09/269,250E

CURRENT FILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 38

SOFTWARE: PatentIn version 3.1

SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-489-760-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-489-760-2
                                                                                                                                                                           ; ORGANISM: Histocompatibility antigen US-09-489-760-2
                                                                                                             Best Loc
Matches
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Matches 9; Conserv
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SEQ ID NO 2
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Best Local
                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                   FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
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100.0%; Pred. No. 4.6e+05;
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                                                                                                           Score 37; DB 2; 1
Pred. No. 4.6e+05;
0; Mismatches 1
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Pred. No. 4.6e+05;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                           Length 9;
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RESULT 7 US-09-217-609A-11

Sequence 11, Application US/09217609A Patent No. 6071733
GENERAL INFORMATION:

APPLICANT: MURAMATSU, Masayoshi

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NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
LENGTH: 9
TYPE: PRT
ORGANISM: Human
FEATURE
                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:

APPLICANT: Rijksumiversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Hunt, Donald F

APPLICANT: Hunt, Donald F

APPLICANT: Hard, Victor H

TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US

CURRENT FILING DATE: 2000-01-21

EARLIER APPLICATION NUMBER: PCT/NL98/00424

EARLIER FILING DATE: 198-07-23

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 1
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PATENT NO. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
                                                                       Matches
                                                                                    Query Match
Best Local S
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LOCATION: (3)
OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE
OTHER INFORMATION: RESIDUE
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                                                                                                                                                                                                                                    ORGANISM: Histocompatibility antigen FEATURE:
                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                                                                                                                                                                                                                        LENGTH: 9
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                1 VLRDDLLEA 9
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                                                                     Similarity
8; Conserv
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VLXDDLLEA
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                                                                     Conservative
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                                                                     Score 36; DB
Pred. No. 4.6e
0; Mismatches
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Pred. No. 4.6e+05;
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                                                                                                                                                                                                                                                                                                                                                                                          US-08-873-235B-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/873,235

FILING DATE: 11-Jun-1997

ATTORNEY/AGENT INFORMATION:

NAMB: TOPPRETTI, Judith L.

REGISTRATION NUMBER: 39,048

REFERENCE/DOCKET NUMBER: 10235/1

TELECOMMUNICATION INFORMATION:

TELEPHONE: 202-429-176

INFORMATION FOR SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                             Sequence 11, Application Patent No. 6174715
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 64.3%;
Best Local Similarity 57.1%;
Matches 4; Conservative
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                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: 1025 Connecticut Avenue, NW - Suite 600
                                                                                                                                                                                   APPLICANT: SHIMIZÚ, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WorddPerfect 6.1 for Window
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WordPerfect 6.1 for Windows CURRENT APPLICATION DATA:
                                                                                                        STREET: 1025 Conr
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 9 amino acids
TYPE: amino acid
                                                          ZIP: 20036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Kenyon & Kenyon
STREET: 1025 Connecticut Avenue, NW - Suite 600
                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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6174715
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OGURA, Kyozo
KOYAMA, Tanetoshi
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Pred. No. 4.6e+05;
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RESULT 10

US-08-556-419-14

; Sequence 14, Application US/08556419C
; Patent No. 6093549
; GENERAL INFORMATION:
APPLICANT: ROSS, Christopher
APPLICANT: Li, Xiao-Jiang
; APPLICANT: Li, Shi-Hua
APPLICANT: Lanahan, Anthony
APPLICANT: Worley, Paul
APPLICANT: Snayder, Solomon
APPLICANT: Snayder, Solomon
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US-09-269-250E-38
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Best Local Similarity
Matches 5; Conserva
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GENERAL INFORMATION:
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1 SEQ ID NO 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Goulmy, E18a
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250B
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 38,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/873,235B
FILING DATE: 11-Jun-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 154441/1996
FILING DATE: 14-Jun-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 202-429-1776
TELEFAX: 202-429-0796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: TOFFENETTI, Judith L. REGISTRATION NUMBER: 39,048 REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
SOFTWARE: WordPer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
INVENTION: Huntingtin-associated protein
                                                                                                                                                                                                                                                                                                                                              h 54.8%; Score 23; DB 2; I
Similarity 100.0%; Pred. No. 4.6e+05;
5; Conservative 0; Mismatches 0;
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nilarity 57.1%;
Conservative
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Pred. No. 4.6e+05;
3; Mismatches 0
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Sequence 4, Application US/09489760

Patent No. 6878375

GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
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US-09-489-760-4
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                                                                                                                                                                                                                                                                                                             RESULT 12
                                                                                                                                                                                                                                                                                            US-08-459-568-20
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Best Local S
Matches 5
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Best Local Similarity 80.0%;
Matches 4; Conservative
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Patent No. 5811304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 4
LENGTH: 9
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CURRENT APPLICATION UNMBER: US/08/556,419C
CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Huang, Shi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KRY: PEPTIDE
LOCATION: (2)..(7)
OTHER INFORMATION: RESIDUES
OTHER INFORMATION: RESIDUES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Histocompatibility antigen FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
                                                                          TITLE OF INVENTION: Retinoblastoma Protein -
TITLE OF INVENTION: Zinc ringer Proteins
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
ADDRESSEB: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite
STREET: 4370 La Joi
CITY: San Diego
STATE: California
CQUNTRY: USA
ZIP: 92122
                                                                                                                                                                                                                                                                                                                                                                                                                                                     y Match 52.4%; Score 22; DB 2;
Local Similarity 55.6%; Pred. No. 4.6e+05;
hes 5; Conservative 0; Mismatches 4
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                                                                                                                                                             Retinoblastoma Protein - Interacting Zinc Finger Proteins

    Mismatches

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Pred. No. 4.6e+05;
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US-08-399-411-20
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Patent No. 5831008
GENERAL INFORMATION:
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Best Local :
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FILING DATE: 02-UN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-WAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 915
REFERENCE/DOCKET NUMBER: 917
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
                                                                                                              APPLICATION UNMER: US/08/399,411
FILING DATE: 06-MAR-1995
CLASSIFICATION: 530
ATTORNEY/ACENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCS/DOCKET NUMBER: P-LJ 1264
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                        TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Retinoblastoma Protein - TITLE OF INVENTION: Zinc Finger Proteins NUMBER OF SEQUENCES: 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   / Match 50.0%;
Local Similarity 80.0%;
les 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U
ZIP: 92122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
TYPE: amino acid TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: San Diego
                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 DDLLE 8
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                                     6 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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Pred. No. 4.6e+05;
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                                                                                                                                                                                                                                                                                                                            Version
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RESULT 15
US-09-586-472-20
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SOFEMARES: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/516,859A
PILING DATE: 18-AUG-1995
CLASSIFICATION: 514
PRIOR APPLICATION: DATA:
APPLICATION NUMBER: US 08/399,411
PILING DATE: 06-MAR-1995
PRIOR APPLICATION NUMBER: US 08/292,663
FILING DATE: 18-AUG-1994
APPLICATION NUMBER: US 08/292,663
FILING DATE: 18-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REFERENCE/DOCKET NUMBER: P-LJ 1776
REFERENCE/DOCKET NUMBER: P-LJ 1776
TELECOMUNICATION: UFFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9999
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                 ; TYPE: amino acid
; TOPOLOGY: linear
US-08-516-859A-20
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                                   Sequence 20, Application US/0
Patent No. 6323335
GENERAL INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: R
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                        Query Match 50.0%;
Best Local Similarity 80.0%;
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION: APPLICANT: Huang,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
ADDRESSEE: Campbell village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: C
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NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                         4 DDLLE 8
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4; Conserv
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                                                                                                               Application US/09586472
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                                                                                                                                                                                                                                                                        Score 21; DB 2; Le Pred. No. 4.6e+05; 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 21; DB 1;
Pred. No. 4.6e+05;
1; Mismatches 0
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Search completed: April Job time: 27.6667 secs
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                                                                                                                                        Query Match
Best Local Similarity 80.0
                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acid
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTEE: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/586,472
FILING DATE: 01-Jun-2000
CLASSIFICATION DATA:
APPLICATION NUMBER: US 09/528,706
FILING DATE: 10-Jun-2000
APPLICATION NUMBER: US 08/516,859
FILING DATE: 11-JMAR-2000
APPLICATION NUMBER: US 08/39,411
FILING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/39,411
FILING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/292,683
FILING DATE: 18-AUG-194
ATTORNEY/AGENT IMPORMATION:
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 4130
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                               4 DDLLE 8
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                    6, 2006, 16:27:37
                                                                                                                                                                                           50.0%;
                                                                                                                                                                     Score 21; DB 2; Length 6; Pred. No. 4.6e+05; Indels 1; Mismatches 0; Indels
                                                                                                                                                                         0,
                                                                                                                                                                     Gaps
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Result
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92
        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        seq length: 0 seq length: 9
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Match Length
      Published Applications AA_Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1867569 seqs, 417829326 residues
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Gapop 10.0 ,
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        GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.
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                 US-10-623-176-10
US-17-791-217-5
US-11-0077-740-18
US-10-623-176-42
US-10-623-176-20
US-10-623-176-20
US-10-623-176-19
US-11-007-740-20
US-11-007-740-20
US-11-007-740-29
US-11-007-740-29
US-11-0623-176-49
US-10-623-176-49
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Sequence 10, Appl
Sequence 5, Appli
Sequence 42, Appli
Sequence 46, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 1, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 41, Appl
Sequence 48, Appl
Sequence 41, Appl
Sequence 47, Appl
Sequence 47, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 349, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 26, Appl
Sequence 27, Appl
Sequence 28, Appl
Sequence 29, Appl
Sequence 26, Appl
Sequence 26, Appl
Sequence 27, Appl
Sequence 28, Appl
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RESULT 2 US-10-791-217-5

Sequence 5, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H

Qy 1 Db 1	Query Match Best Local Si Matches 9;	; FAATORE; ; NAME/KEY: ; LOCATION: US-10-623-176-	; FEATURE:	- ⊢	SEQ ID NO 1	; NUMBER OF S ; SOFTWARE: P	; PRIOR APPLI ; PRIOR FILIN			; PRIOR APPLI		ää	; TITLE OF IN ; FILE REFERE	; APPLICANT: ; APPLICANT:	_	US-10-623-176-10; Sequence 10, Ap; Publication No.	RESULT 1		44 21 45 21		42 21			37 21							•
VLRDDLLEA 9 VLRDDLLEA 9	100.0 Similarity 100.0 9; Conservative	SITE (1)(9) 10	E: INFORMATION: Description	Artificial Sequence		SEQ ID NOS: 77 PatentIn Ver. 2.	DATE: 20	1998-0	FILING DATE: 1997-07-23	APPLICATION NUMBER: EP	APPLICATION NUMBER: 09/489,	APPLICATION NUMBER: FILING DATE: 2003-	TITLE OF INVENTION: HA-1 epitope FILE REFERENCE: 2183-6047US	rd,	Goulmy, Els A.J.	pplication . US2004009			50.0	50.0	50.0	50.0	50.0	50.0	50.0	л (л О	50.0	52.4	52.4	54.8	54.8
	0; Sc		Of.	ence		ב	JP 2000-504165 -24	-23	3 7/NT.98/004	97202303.	09/489,760	US/10/623, 07-18	s and	tor H.	.M.	US/10623176 92446A1		ALIGN	US-	-sn	3 US-09-93	SD	Su	SU	SS			US-11-	US-10-	US-10-	3 US-09-83
	ore 42; DB 4; Le ed. No. 1.7e+06; Mismatches 0;		Artificial Sequence:				65	i d	24	0		176	uses thereof					ALIGNMENTS	10-119-536A-109	5-430-565	-09-935-430-219	7-268-9	2-675-6	-09-989-994-211	US-09-989-994-208	0-186-211	US-09-990-186-208	740-	.791-217-4	176-	4-765-132
	Length 9; Indels 0;		ce: HA-1 peptide																Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
	Gaps 0;		ide																e 54, Appı e 109, App	565,	e 219, App	9		212	208,	211,	208,	40,	44	ω	e 132, App

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RESULT 4
US-10-63-176-42
; Sequence 42, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
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APPLICANT: Goulmy, Els A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

APPLICANT: Engelhard, Victor H.

FITTLE OF INVENTION: HA-1 epitopes and uses the FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR FILING DATE: 2000-01-21

PRIOR APPLICATION NUMBER: EP 97202303.0
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Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 1999-05-21
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARD: DEFORTE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
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SEQ ID NO 18
LENGTH: 9
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 9
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: obtained from KIAA0223 partial complementary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                 1 VLRDDLLEA 9
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Sequence 46, Application US/10623176

| Publication No. US20040092446A1
| CENERAL INFORMATION:
| APPLICANT: Hunt, Donald F.
| APPLICANT: Hunt, Donald F.
| APPLICANT: Hunt, Donald F.
| APPLICANT: Engelhard, Victor H.
| APPLICANT: Engelhard, Victor H.
| APPLICANT: Engelhard, Victor H.
| CURRENT PILING DATE: 2183-6047US
| CURRENT APPLICATION NUMBER: US/10/623,176
| CURRENT FILING DATE: 2000-01-21
| PRIOR APPLICATION NUMBER: EP 97202303.0
| PRIOR APPLICATION NUMBER: EP 97202303.0
| PRIOR APPLICATION NUMBER: EP 97202303.0
| PRIOR APPLICATION NUMBER: BP 97202303.0
| PRIOR APPLICATION NUMBER: J9 2000-504165
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US-10-623-176-46
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; LOCATION: (1)..(9)
US-10-623-176-42
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-46
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SEQ ID NO 42
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PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR PILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                 LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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                                                                                                                                                                                                                                                                                        FEATURE:
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                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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                                                                                                         Local
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                                                                                                       90.5%; Score 38; DB 4; 100.0%; Pred. No. 1.7e+
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US-10-623-176-2 ; Sequence 2, Application US/10623176 ; Publication No. US20040092446A1

RESULT 6

GENERAL INFORMATION:

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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-2
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US-10-791-217-2
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: DT 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-1-24
                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/10791217 Publication No. US20040191268A1
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Best Local Similarity
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SEQ ID NO 2
                                                                                                                                                                                                                                                                                                          CURRENT PPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR PILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR PILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Goulmy, APPLICANT: Hunt, I APPLICANT: Engelha
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses
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TYPE: PRT
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                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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                    VLRDDLLEA 9
VLHDDLLEA 9
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                                                                             Conservative
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                                                                       Score 37; DB 4; Length 9; Pred. No. 1.7e+06; 0; Mismatches 1; Indels
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Pred. No. 1.7e+06;
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RESULT 10
US-10-623-176-1
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US-11-007-740-20
US-11-007-740
Sequence 20, Application US/11007740
Fublication No. US20050233350A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                   , OTHER INFORMATION: Description of Artificial Sequence: Exon US-11-007-740-20
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SEQ ID NO 20
LENGTH: 9
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APPLICANT: Goulmy, Elea A.J.M.
APPLICANT: Goulmy, Elea A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunc TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: PCT/NLO2/00791
PRIOR APPLICATION NUMBER: EPT/NLO2/00791
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.1
Sequence 1, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 8; Conserv
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Best Local Similarity
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
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APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.

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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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Best Local Similarity 88.
Matches 8; Conservative
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Publication No. US20040191268A1
GENERAL INFORMATION:
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                                                                                            Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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NAME/KRY: MISC FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide OTHER INFORMATION: wherein X can be R or H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: SITE
LOCATION: (1)..
                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial sequence
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                                                                        Local Similarity nes 8; Conserv
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                                      1 VLRDDLLEA 9
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Pred. No. 1.7e+06;
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                                                                        Score 36; DB 4;
Pred. No. 1.7e+06;
0; Mismatches 1
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                                                                              Indels
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; OTHER INFORMATION: E ; FEATURE: ; NAME/KEY: SITE ; LOCATION: (1)..(9) US-10-623-176-48
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; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (3)...(3)
; OTHER INFORMATION: Xaa represents a histidine (H)
US-11-007-740-29
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP7/012303.0
PRIOR APPLICATION NUMBER: EPT/NL98/00424
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 2000-01-24
NUMBER: OF 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER: DESCRIPTION UMBER: PT/NL98/00424
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US-11-007-740-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1 SEQ ID NO 29
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Best Local (
                                                            Best Loc
Matches
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SEQ ID NO 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 48, Application US/10623176
Publication No. US20040092446A1
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CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: GOULMY, E18A
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
                                                                                                     Query Match
                                                                                                                                                                                                                                                  TYPE: PT
TYPE: PT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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Local Similarity 88.9%;
                                                          Local Similarity
les 7; Conserv
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3 RDDLLEA 9
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                                                                Conservative
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                                                              81.0%; Score 34; DB
100.0%; Pred. No. 1.7
tive 0; Mismatches
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Pred. No. 1.7e+06;
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US-10-623-176-23

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Sequence 23, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
FILL OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT PILING DATE: 2003-07-18
FRICA APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-24
NUMBER OF SEQ ID NOS: 77
SOOTMARE: PATHONISM: ANTIGINAL NOTE: 2.1
SEQ ID NO 23
LENGTH: 9
TYPE: PRI
OCCANISM: ANTIGINAL NOTE: 2.1
                                                                                                               PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR PPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTMARE: PATENTIN Ver: 2.1
SEQ ID NO 41
PRIOR TILING DATE: 2000-01-24
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US-10-623-176-41
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-23
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Best Local Similarity lou.
"Arches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 41, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR PILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                ORGANISM: Artificial Sequence
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100.0%; Pred. No. 1.7e+06;
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; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-41

Query Match
Best Local Similarity 87.5%; Pred. No. 1.7e+06;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps
Qy 1 VLRDDLLE 8
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Db 2 VLHDDLLE 9
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0

Search completed: April 6, 2006, 16:50:29 Job time : 90.3333 secs THIS PAGE BLANK (USPTO)

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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         Pred. No. is the number of results predicted a score greater than or equal to the score of the and is derived by analysis of the total score
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         length: 0 length: 9
       Match
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/SIDS5/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US11_NEW_PUB.pep:*
/SIDS5/ptodata/2/pubpaa/US11_NEW_PUB.pep:*
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US-11-010-748A-12
US-11-010-748A-11
US-11-010-748A-11
US-11-010-748A-11
US-11-026-706-52
US-11-225-686-208
US-11-225-686-211
US-11-225-686-211
US-11-225-686-211
US-11-225-099-216
US-11-202-009-216
US-11-202-009-216
US-11-202-009-216
US-11-202-009-216
US-11-202-009-216
US-11-209-143-168
US-11-129-143-173
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                       Sequence
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  12, Appl
17, Appl
208, Appl
208, Appl
216, App
216, Appl
216, Appl
216, Appl
216, Appl
216, Appl
168, Appl
171, Appl
172, Appl
173, Appl
174, Appl
175, Appl
176, Appl
177, Appl
177, Appl
178, Appl
179, Appl
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ALTCHMENTS	US-11-045-024-5716	US-11-045-024-5495	US-11-045-024-3698	US-11-045-024-3138	US-11-045-024-575	US-11-202-009-3941	US-11-202-009-3932	US-11-202-009-3916	US-11-202-009-3915	US-11-202-009-3914	US-11-202-009-3877	US-11-202-009-2954	US-11-225-686-3941	US-11-225-686-3932	US-11-225-686-3916	US-11-225-686-3915	US-11-225-686-3914	US-11-225-686-3877	US-11-225-686-2954	US-10-927-634-63
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	5716, 1	5495, 1	3698, 1	3138, 1	575, A	3941, 1	3932, 1		3915, 3	3914,	3877,	2954,	3941,			3915,	3914,	3877,	54	63, Appl
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TYPE: PRT
ORADISM: Homo sapiens
FEATURE:
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)...(9)
TOTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by
TOTHER INFORMATION: ell epitopes
US-11-010-748A-12
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US-11-010-748A-12
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Publication No. US20050244421A1
GENERAL INFORMATION:
APPLICANT: Merck Patent GmbH
APPLICANT: STRITTMATTER, Wolfgang
APPLICANT: SCHARM, Heidrun
APPLICANT: SCHARM, Burkhard
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE
                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.1
SEQ ID NO 12
LENGTH: 9
                                                                                                                   Query Match
                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: MER-136
CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
PRIOR APPLICATION NUMBER: PCT/EP03/06251
PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2002-0
1 VLRDDLLEA 9
                                                                       Similarity 100.0%; Score 42; Similarity 100.0%; Pred. No. 9; Conservative n. ....
                                                                                                                                                                                                                                                                                                                                                                                                2002-06-13
                                                                              Mismatches
                                                                                                1.4e+05;
                                                                                                                       DB 7;
                                                                                                                   Length 9;
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RESULT 2 US-11-010-748A-11

Sequence 11, Application US/11010748A
Publication No. US20050244421A1
GENERAL INFORMATION:
APPLICANT: Merck Patent GmbH
APPLICANT: STRITTWATTER, Wolfgang
APPLICANT: MOLL, Heidrun
APPLICANT: SCHARM, Burkhard

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RESULT, 4
US-11-096-706-52
; Sequence 52, Application US/11096706
; Publication No. US20050245476A1
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Matches
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                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.2 SEQ ID NO 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ZHOU, Hongxing
APPLICANT: ZHOU, Chen
APPLICANT: COSMAN, David J.
APPLICANT: CORTER, Paul
APPLICANT: CARTER, Paul
TITLE OF INVENTION: METHODS OF IDENTIFYING FUNCTIONAL ANTIBODIES
FILE REFERENCE: A-890A
CURRENT APPLICATION NUMBER: US/11/019,027
CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: 60/605,902
PRIOR PILING DATE: 2004-08-31
PRIOR APPLICATION NUMBER: 60/531,714
PRIOR APPLICATION NUMBER: 60/531,714
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: YAN, APPLICANT: SHE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE
                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2003-
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (1). (9)
OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T
OTHER INFORMATION: ell epitopes
                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial sequence
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                                                                                                                                                                                                                                                                                          OTHER INFORMATION: ER localization signal
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100.0%; Pred. No. 1.4e+05;
tive 0; Mismatches 0;
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US-11-225-686-208
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LENGTH: 7
TYPE: PRT
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APPLICANT: Sangamo Biosciences, Inc.
APPLICANT: Sangamo Biosciences, Inc.
APPLICANT: Collingwood, Trevor
IITLE OF INVENTION: Treatment of Neuropathic Pain with Zinc Finger Proteins
FILE REFERENCE: 019496-008220US
CURRENT APPLICATION NUMBER: US/11/096,706
CURRENT PILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 60/560,535
PRIOR APPLICATION NUMBER: US 60/560,535
PRIOR APPLICATION NUMBER: US 60/576,757
PRIOR PILING DATE: 2004-06-02
NUMBER OF SEQ ID NOS: 227
RUMBER OF SEQ ID NOS: 227
NUMBER OF SEQ ID NOS: 227
NUMBER OF SEQ ID NOS: 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 208, Application US/11225686

Publication No. US20060019349A1

GENERAL INFORMATION:

APPLICANT: LIU, Qiang

TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE

TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS

FILE REFERENCE: 8325-0011.20 / $11-US2

CURRENT APPLICATION NUMBER: US/11/225,686

CURRENT FILING DATE: 2005-09-12
Sequence 211, Application US/11225686

Publication No. US20060019349A1

GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REPERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/225,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 208
LENGTH: 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2005-09-12
PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Chemically synthesized peptide (zinc finger protein recognition OTHER INFORMATION: region)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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| |||:
1 RSDLLQ 6
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Pred. No.
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US-11-202-009-208
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / $11-U$2
CURRENT APPLICATION NUMBER: U$/11/225,686
CURRENT FILING DATE: 2005-09-12
PRIOR APPLICATION NUMBER: U$/09/989,994
PRIOR APPLICATION NUMBER: U$/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 216
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Best Local Similarity 65...
"Thes 4; Conservative
                                                      APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT PLILOR ON NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/99/989,994
PRIOR RILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 208
                                                                                                                                                                                                                                                                              Sequence 208, Application US/11202009 Publication No. US20060024726A1 GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 211
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Best Local Similarity
Matches 4; Conser
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: example ZFP
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ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Artificial Sequence
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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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US-11-202-009-216
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PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 211
                                                                                                                                                                                                      TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS FILE REFERENCE: 8325-0011.20 / $11-US2 CURRENT APPLICATION NUMBER: US/11/202,009 CURRENT FILING DATE: 2005-08-11 PRIOR APPLICATION NUMBER: US/09/989,994 PRIOR PILING DATE: 2001-11-20 NUMBER OF SEQ ID NOS: 4085 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 216
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Publication No. U320060024726A1
GENERAL INFORMATION:
APPLICANT: LIU, Olang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLECTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / $11-US2
CURRENT APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
                                Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
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TYPE: PRT
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ORGANISM: Artificial Sequence
PEATURE:
                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: example
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3 RDDLLE 8
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                                Score 21; DB 7;
Pred. No. 1.4e+05;
1; Mismatches 1
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Pred. No. 1.4e+05;
1; Mismatches 1
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Pred. No.
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APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
APPLICANT: LEVIN, ELANA
APPLICANT: CANALITA-BID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 00/935,430
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR PRIOR DATE: 2001-08-22
PRIOR PRIOR DATE: 2001-08-22
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PRIOR FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR PILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 219
LENGTH: 9
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Best Local Similarity
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TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
CURRENT FILING DATE: 2004-11-15
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APPLICANT: HUBERT, RENE
APPLICANT: RAITANO, ARTHUR
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RAITANO, ARTHUR
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Pred. No. 1.4e+05;
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 565
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 169
LENGTH: 5
TYPE: PRT
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Publication No. US20050266518A1
GENERAL INFORMATION:
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SEQ ID NO 168
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APPLICANT: BERRY, Alan
APPLICANT: BERTZEL, Werner
APPLICANT: HUMBELIN, Markus
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Best Local Similarity 37.1
Matches 3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: LOPEZ-ULÍBARRI, RUAL
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEBY, Alexei A.
RITILE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
RUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
                                                                                                                                      APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Weriner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: WAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 5
TYPE: PRT
ORGANISM: Bradyrhizobium japonicum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 VLRDDLLE 8 ::: ||:| 1 LMKSDLVE 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 DDLLE 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  47.6%; Score 20; DB 7;
60.0%; Pred. No. 1.4e+05;
tive 2; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 21; DB 6;
Pred. No. 1.4e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 5;
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US-11-129-143-170

US-11-129-143-170

Sequence 170, Application US/11129143

Publication No. US20050266518A1

GENERAL INFORMATION:
APPLICANT: BERTZY, Alan
APPLICANT: HUMBELIN, Markus
APPLICANT: HUMBELIN, Markus
APPLICANT: VELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION VUMBER: US/11/129,143
CURRENT PILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
LENGTH: 5
TYPE: PRT
ORGANISM: Bacillus stearothermophilus
US-11-129-143-170
Search completed: April 6, 2006, 16:51:12 Job time : 12.3333 secs
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US-11-129-143-169
                                                                                                                                                                        Query Match 47.6%; Score 20; DB 7; Length 5; Best Local Similarity 60.0%; Pred. No. 1.4e+05; Matches 3; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 47.6%; Score 20; DB 7; Length 5; Best Local Similarity 60.0%; Pred. No. 1.4e+05; Matches 3; Conservative 2; Mismatches 0; Indels
                                                                                     4 DDLLE 8 | ||:|:
1 DDILD 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 DDLLE 8 | | :|:
1 DDILD 5
                                                                                                                                                                          0; Indels
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